Copyright

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Minimum DB
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1: geneseqp1980s:*
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3: geneseqp2000s:*
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and is derived is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	σı	4	ω	N	щ	NO.	Result
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Aam50103 Equine Ig	Aab06206 Immunogen	Aaw24102 Canine im	Abp96589 Rat IgE h	Abg74785 Human CH2	Rat	Rat	Cat	Adg73251 Cat parti	Abu09336 Feline Ig	Fel	_	Adg73237 Cat immun	Immu	Abp96583 Dog IgE h	Aar97753 Canine Ig	Aaw23067 Canine Ig	Abg74782 Human CH2	Human		Aay79995 Dog immun	Parti	_	Human	Abg74772 Canine Ig	Description	

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65.4	65.4	66.5	66.5	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	68.8	69.1	70.2	70.2	70.2	72.1	12.1
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ALIGNMENTS

RESULT 1 ABG74772 05-JUN-2003 (first entry) ABG74772 standard; protein; 114 ABG74772; B

Canine IgB

CH3 domain.

CH3 domain; IgB; antigen; non-anaphylactic; anti-IgB; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergy; thinitis; gastrointestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis; gene therapy; dog.

Canis familiaris.

EP1262491-A2

04-DEC-2002.

22-MAY-2002; 2002EP-00253606

22-MAY-2001; 2001US-0292638P.

(PFIZ) PFIZER PROD INC.

Brown TM, Morsey MA;

N-PSDB; WPI; 2003-122561/12. ACA55170.

Novel isolated antigenic peptide comprising amino acid residues of CH3 domain of IgE molecule from first species and a second unrelated species, induces non-anaphylactic anti-IgE immune response in animal.

Claim 15; Page 28; 50pp; English.

This invention describes a novel antigenic peptide comprising amino acid residues of an IgE CH3 domain from a first species (ADE1) and amino acid residues of an IgE CH3 domain of a second unrelated species (ADE2), where ADE1 is conserved in the IgE CH3 domain of the second species and ADE2 is not conserved in the IgE CH3 domain of the first species. The novel antigenic peptide induces a non-anaphylactic anti-IgE immune response in an animal. The invention also discloses the polymucleotide sequence encoding the antigenic peptide and an antigenic fusion protein comprising the antigenic peptide of the invention and a heterologous protein

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RRESULT 2
ABG74774
ID ABG74774
AC ABG7
XX ABG7
XX ABG7
XX ABG7
XX ABG7
XX CH3
XX CH3
XX CCA1
XX Germ
XX Germ
XX Germ
XX Germ
XX Germ
XX Germ
XX Homo
OS Chim
XX EP12
XX EP12
XX EP12
XX EP12
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Best Local S
Matches 25
                           Claim
                                                   Novel isolated antigenic peptide comprising amino acid residues of CH3 domain of IgE molecule from first species and a second unrelated specinduces non-anaphylactic anti-IgE immune response in animal.
                                                                                                                                                                                              N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                 22-MAY-2001; 2001US-0292638P.
                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-2002; 2002EP-00253606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CH3 domain; IgB; antigen; non-anaphylactic; anti-IgB; fusion protein; dermatcological; antiinflammatory; ophthalmological; allergy; asthma; allergic rhinitis; gastrointestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis;
                                                                                                                                                                                                                                                                                                                                   (PFIZ ) PFIZER PROD
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05-JUN-2003
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                                                                                                                                                                                        2003-122561/12.
DB; ACA55172.
         2; Page 29-30; 50pp; English
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                                                                                                                                                                                                                                                                             Morsey MA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
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RESULT 3
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Matches 25
                                                      Brown TM,
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                                                                                                                                                           22-MAY-2001; 2001US-0292638P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CH3 domain; IgE; antigen; non-anaphylactic; anti-IgE; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergic rhinitis; gastrointestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis;
                                                                                                                                                                                                                           22-MAY-2002; 2002EP-00253606.
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                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
Canis familiaris.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy; human; dog.
                                                                                                          ) PFIZER PROD INC.
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25; Conserva
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                                                Morsey MA;
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Pred. No. 6.
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Claim 2; Page 29; 50pp; English.

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ID AAW
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PD 30--
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     WPI; 1997-389423/36
                                                                                                                                                                                                                                                                                                                                  22-DEC-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunoglobulin E; IgE; anti-canine IgE antibody; allergy; canine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Partial canine immunoglobin E protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW24097 standard; peptide; 124
                                                                                                                 (HITB ) HITACHI CHEM CO LTD
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antigenic peptides (I). (I) have anti-allergic, anti-anaphylactic and anti-asthmatic properties. (I) induces polyclonal antibodies specific a target effector site on the epsilon-heavy chain of IgS, and so preventing triggering and activation of mast cells and basophils and downregulation of IgS synthesis. Conjugates, or fusion peptides, containing (I) are used for active immunisation against IgS-mediated allergies, e.g. food allergies, asthma, anaphylaxis, or flea-allergy dermatitis. Nucleic acids that encode these compounds are useful for recombinant production of corresponding peptides or in DNA vaccines. Conjugates of (I) that include a promiscuous T helper cell epitope (functional in genetically diverse subjects), in addition to a B cell target epitope.

The present invention describes immunoglobulin E (IgE)-CH3 domain

for

Example

1; Page 66-68; 155pp; English

Query Match Best Local S Matches 25

al Similarity 25; Conserv

100.0%; Score 136; DB 6; llarity 100.0%; Pred. No. 6.7e-13; Conservative 0; Mismatches 0;

Length 117;

<u>.</u>.

Sequence 117 AA,

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This invention describes a novel antigenic peptide comprising amino acid considered of an IgE CH3 domain from a first species (ADE1) and amino acid considered of an IgE CH3 domain of a second unrelated species (ADE2), where CC ADE1 is conserved in the IgE CH3 domain of the second species and ADE2 is conserved in the IgE CH3 domain of the first species. The novel control of the invention also discloses the polynucleotide sequence control of the invention also discloses the polynucleotide sequence control of the invention and an antigenic fusion protein comprising the antigenic peptide of the invention and a heterologous protein comprising that does not cause anaphylaxis when administered to an animal. The conducts of the invention have dermatological, antiinflammatory and cophthalmological activity. The antigenic peptide described is capable of products of the invention have dermatological, antiinflammatory and cophthalmological activity. The antigenic peptide described is capable of compressioned to a medicament for treating or preventing IgE-mediated allergic disorders including asthma, allergic rhinitis, gastrointestinal allergies such as food allergies, eosinophilia, conjunctivitis, glomerular nephritis, flea conjunctivitis, granting ige-mediated allergic disorders by gene therapy. Antigenic peptides comprising conserved animo acid residues of the CH3 domain of an IgE molecule from one species considered by variable amino acid residues of the CH3 domain of an IgE conjunction and antibodies are capable of inducing a high citre of anti-IgE antibodies when administered to an animal without conserved in designing the constructs described in the disclosure of the conserved in the disclosure of the constructs described in the disclosure of the constructs described in the disclosure of the conserved in the disclosure of the conserved conserved animal without conserved in the disclosure of the conserved cons
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunogenic; immunostimulatory; cas
antibody; allergy; allergic disease
anti-anaphylactic; anti-asthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This is a partial canine immunoglobulin E (IgE) protein. Peptide fragments (AAW24098-106) containing at least five continuous amino acids of this sequence are used for the preparation of anti-canine IgE antibody. The anti-canine IgE antibody can be used for the diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY79995 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  canine allergies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wang CY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunoglobulin E; IgE; epsilon heavy chain; antigenic; antigen;
immunogenic; immunostimulatory; carrier protein; helper T cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dog immunoglobulin B epsilon heavy chain SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UNBI-) UNITED BIOMEDICAL INC
                                                                                                                                                                                                                                                                                                                                                                        antigenic peptide from the CH3 domain immunization against allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 CRVTHPHLPKDIVRSIAKAPGKRAP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 8; 12pp;
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100.0%; Pred. No. 7.2e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease; immunisation; anti-allergic
nmatic; asthma; anaphylaxis; dermatiti
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This invention describes a novel antigenic peptide comprising amino acid CC residues of an IGE CH3 domain from a first species (ADE1) and amino acid CC residues of an IGE CH3 domain of a second unrelated species (ADE2), where CC ADE1 is conserved in the IGE CH3 domain of the first species and ADE2 is CC not conserved in the IGE CH3 domain of the first species. The novel CC antigenic peptide induces a non-anaphylactic anti-IGE immune response in CC an animal. The invention also discloses the polynucleotide sequence CC the antigenic peptide of the invention and a heterologous protein comprising CC carrier, where the fusion protein induces an anti-IGE immune response that does not cause anaphylaxis when administered to an animal. The CC ophthalmological activity. The antigenic peptide described is capable of products of the invention have dermatological, antiinflammatory and CC ophthalmological activity. The antigenic peptide described is capable of preventing IGE from binding to high affinity receptors on mast cells and basophils. The products of the invention are useful in the manufacture of inclinding asthma allergic disorders.
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Best Local (
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                                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CH3 domain; IgB; antigen; non-anaphylactic; anti-IgB; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergic rhinitis; gastrointestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PFIZ ) PFIZER PROD INC
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Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        constraints (disulfide bridge) to stabilise conformational features and maximize cross-reactivity to the natural target. They induce safe (non-anaphylactogenic) antibodies. AAY79994 to AAY80084 represent amino acid sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human IgB CH2-canine
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                                                                                                                                                                                                                                                                                                                 3; Page 34-36; 50pp; English.
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     asthma,
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Pred. No. 2e-12;
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gastrointestinal allergies
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RESULT 7
ABG747933
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             This invention describes a novel antigenic peptide comprising amino acid residues of an IgE CH3 domain from a first species (ADE1) and amino acid residues of an IgE CH3 domain of a second unrelated species (ADE2), where ADE1 is conserved in the IgE CH3 domain of the second species and ADE2 is not conserved in the IgE CH3 domain of the first species. The novel antigenic peptide induces a non-anaphylactic anti-IgE immune response in an animal. The invention also discloses the polymucleotide sequence encoding the antigenic peptide and an antigenic fusion protein comprising the antigenic peptide of the invention and a heterologous protein
                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated antigenic peptide comprising amino acid residues of domain of IgB molecule from first species and a second unrelated spinduces non-anaphylactic anti-IgB immune response in animal.
                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 37-39; 50pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; ACA55181.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human CH2-human/canine CH3-human CH4 IgE-3 fusion protein.
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s familiaris.
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ilarity 100.0%;
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Pred. No. 2.3e-12;
); Mismatches 0;
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fusion protein

induces an anti-IgE

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CC that does not cause anaphylaxis when administered to an animal. The CC products of the invention have dermatological, antiinflammatory and CC ophthalmological activity. The antigenic peptide described is capable of preventing IgE from binding to high affinity receptors on mast cells and CC preventing IgE from binding to high affinity receptors on mast cells and CC basophils. The products of the invention are useful in the manufacture of CC anedicament for treating or preventing IgE-mediated allergic disorders including asthma, allergic rhinitis, gastrointestinal allergics such as CC food allergies, eosinophilia, conjunctivitis, glomerular nephritis, flea CC allergies or atopic dermatitis, in an animal, e.g. human or dog. The CC disorders, by gene therapy. Antigenic peptides comprising conserved amino CC disorders, by gene therapy. Antigenic peptides comprising conserved amino CC acid residues of the CH3 domain of an IgE molecule from one species CC flanked by variable amino acid residues of the CH3 domain of an IgE CC molecule from a second unrelated species are capable of inducing a high CC causing anaphylaxis. ABG74772-ABG74785 represents polypeptide sequences CC invention
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CH3 domain, IgE, antigen; non-anaphylactic; anti-IgE; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergic thinitis; gastrointestins allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis; gene therapy; human; canine; dog.
Novel isolated antigenic peptide comprising amino acid residues of domain of IgE molecule from first species and a second unrelated synduces non-anaphylactic anti-IgE immune response in animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAY-2002; 2002EP-00253606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABG74782 standard; protein; 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-MAY-2001; 2001US-0292638P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2002.
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Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CH2-human/canine CH3-human CH4 IgE-2 fusion protein.
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                                                                                                                                                                                                                                                                                                                                                                                 (PFIZ ) PFIZER PROD INC
                                                                                                                                                                          2003-122561/12.
DB; ACA55180.
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                                                                                                                                                                                                                                                                                                 Morsey MA;
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This invention describes a novel antigenic peptide comprising amino residues of an IgE CH3 domain from a first species (ADE1) and amino residues of an IgE CH3 domain of a second unrelated species (ADE2),

acid acid where

WO9730156-A2

Page 36-37; 50pp;

English.

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RRSULT 9
AAW230
AAW27
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AAW2
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AAW2
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IGB;
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IGG;
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Best Local :
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                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canine IgE heavy chain constant region (exon 1-4 product).
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                                                                                                                                 Misc-difference
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hypersensitivity; therapy; dog; antisense; immunomodulation.
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176
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175
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-FEB-1997;
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                                                        N-PSDB;
                                                                                          Hollis GF,
                                                                                                                                     09-NOV-1994;
09-NOV-1994;
                                                                                                                                                                       03-NOV-1995;
                                                                                                                                                                                                                                          Canis familiaris.
                                                                                                                                                                                                                                                              IgE; immunoglobulin; antibody; vaccine; allergy; hypersensitivity.
                                                                                                                                                                                                                                                                                      Canine
                                                                                                                                                                                                                                                                                                            28-AUG-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 417 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      products
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IDEX-) IDEXX LAB INC.
                                                                    WPI; 1996-277321/28.
 Claim 11; Page 29-30; 49pp; English
                      New DNA encoding canine IgE and IgA -
therapy, assays, drug screening, etc.
                                                                                                                                                                                             23-MAY-1996.
                                                                                                               (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                               289
                                                                                                                                                                                                                                                                                     IgE.
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                                                          AAT29824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    canine IgE heavy chain constant region DNA - useful to develop for treatment of canine allergies and for immunomodulation in
                                                                                                                                                                                                                                                                                                                                                                                                              ĊŔVTHPHĹPKĎÍVŘSIAKAPGKRAP 313
                                                                                                                                                                                                                                                                                                                                                                                                                                  CRVTHPHLPKDIVRSIAKAPGKRAP 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Harris RA,
                                                                                          Patel MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                     94US-00336583.
94US-00336891.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 136; DB 2;
100.0%; Pred. No. 2.8e-12;
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                                    useful
                                   'n
                                   vaccines, anti-sense
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RESULT 11
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       Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The canine IgB amino acid sequence (AAR97753) was deduced from an isolated gene (AAR29824) obtd. from a canine liver DNA library. The cloning of the IgB gene allows prodn. of large quantities of recombinant IgB using bacterial, yeast, mammalian, insect or viral systems. The IgB can be used in drug development (e.g. small molecule screening, assay development and anti-IgB antibody generation). Fragments of IgB can be used in vaccines or to prevent IgB-mediated hypersensitivity. The new sequence information permits targeted modulation of IgB-mediated immune
The present invention describes a method (M1) for identifying peptides that induce cytotoxic T-lymphocyte (CTL) response against immunoglobulin E (IgE), comprising providing a test peptide (T) suspected of being able to bind to major histocompatibility complex (MHC) class I molecule, and evaluating (T) for ability to elicit in a mammal a CTL response to naturally processed and presented IgE peptides, where a peptide that induces such a response is identified. Also described are compositions: (C1) comprising at least one immunogenic peptide (I) identified by (M1); (C2) comprising at least one isolated polynucleotide encoding (I); and (C3) comprising antigen-presenting cells that recognise at least one (I).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunoglobulin E; vaccine; IgE; cytotoxic T lymphocyte response; immune response; major histocompatibility complex; MHC; immunogenic; antiallergic; antiasthmatic; immunosupressive; vasotropic; cytostatic; dermatological; antiinflammatory; IgE-mediated condition; food allergy; atopic hypersensitivity condition; allergic rhinitis; allergic asthma; atopic dermatitis; non-atopic hypersensitivity condition; anaphylaxis;
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                                                                                                                                                                                                                                                                                                                                                                         Chen
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                                                                                                                                                                                                                                                         Identifying peptides that induce cytotoxic T-lymphocyte, CTL response against IgE, by identifying peptide eliciting CTL response to IgE peptides naturally presented by major histocompatibility complex class I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dog IgE heavy chain amino acid sequence SEQ
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                                                                                                                                                                                                                                               protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-AUG-2002; 2002WO-US026986.
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                                                                                                                                                                                                         Example 7; Page 152-154; 187pp; English.
                                                                                                                                                                                                                                                                                                                                       WPI; 2003-268242/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-AUG-2001; 2001US-0312120P
                                                                                                                                                                                                                                                                                                                                                                         SA,
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                                                                                                                                                                                                                                                                                                                                                                         Yang Y,
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                                                                                                                                                                                                                                                                                                                                                                           Barankiewicz T,
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Pred. No. 2.9e-12;
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RESULT 12
AAB06208
ID AAB06
XX Dog;
XX Didel
OS Chime
XX AAB06
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XX Dog;
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OS Chime
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Best Local :
The present sequence is an immunogenic peptide consisting of the heavy chain constant regions 2 and 4 of the opossum IgB and the heavy chain constant region 3 from the dog. It was shown to cause a stronger polyclonal anti-self IgB response than peptides consisting of the same regions from one mammal. Immunogenic peptides, particularly those consisting of different heavy chain constant regions, can be used for vaccination in humans, against bacterial and viral infections and allergies, such as asthma, fur, pollen and food allergies and eczema. (Updated on 12-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             where C1-3 are able to bind to at least one MHC class I molecule and to elicit in a mammal a CTL response to naturally processed and presented IGE peptides. C1-3 have antiallergic, antiasthmatic, immunosuppressive, vasotropic, dermatological, antiinflammatory and cytostatic activities, and can be used as inducers of a CTL response against IGE, and in vaccines. C1-3 can be used for modulating an IGE-mediated condition such as IGE-mediated atopic hypersensitivity condition. IGE-mediated non-atopic hypersensitivity condition, IGE myslomal. Preferably, C1-3 are useful for treating atopic hypersensitivity conditions (such as allergic rhinitis, allergic asthma, food allergies, or atopic dermatitis), non-atopic hypersensitivity conditions (such as anaphylaxis, and urticaria hives). The present sequence represents an IGE heavy chain amino acid sequence, which is given in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-NOV-1998;
22-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunogenic polypeptides useful for immunoglobulin E in mammals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-365342/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunogenic peptide consisting of opossum CH2, dog CH3 and opossum CH4.
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22-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RESI-) RESISTENTIA PHARM AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  opossum; immunoglobulin E; IgE; vaccination; infection; allergy; na; eczema; immunogenic peptide.
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                                                                                                                                                                                                                                                                                                                                                            Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .Conservative
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(first entry)
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99US-00401636
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                                                                                                                                                                                                                                                                                                                                                            2; 50pp;
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Pred. No. 2.9e-12;
D; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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RESULT 14
ABP96580
ID ABP96
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AC ABP96
XX

ABP96580 standard;

protein;

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ABP96580;

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RESULT 13
ADG73237
ID ADG73
XX ADG74
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                                                                                              Query Match
Best Local S
Matches 21
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                                                                                                                                                                                                                                           The invention describes an isolated nucleic acid molecule (I) encoding a portion of a feline IGE heavy chain protein. The methods and composition of the present invention are useful for eliciting feline immune response for and/or treating IGE-mediated responses, such as allergies. This is the amino acid sequence of a cat immunoglobulin E (IGE) constant region.
                                                                                                                                                                                               Sequence 431
                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid molecule encoding a portion of a feline IgE heavy chain protein, useful for treating and/or eliciting feline immune responses for IGE-mediated responses, such as allergies.
                                                                                                                                                                                                                                                                                                                                                                                                   Claim 12,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-010802/01.
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07-JAN-2000; 2000US-00479614
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cat immunoglobulin B (IgE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MCCA/) MCCALL C. (WEBE/) WEBER B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
299
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                                                                                                21;
                                            1 CRVTHPHLPKDIVRSIAKAPGKRAP
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                                                                                                                     Similarity
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CKVTHPDLPKDIVRSIAKAPGRRFP
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                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weber B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                   84.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                   44pp; English.
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Pred.
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Pred. No. 5.7e-10;
0; Mismatches 1
                                                                                                Mismatches
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                                                                                                                   115;
No. 5.
323
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                                                                                   DB 8; 5.3e-09; 2;
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                                                                                                                                          Length 431;
                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                       compositions
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                                                                                                                                                                                                                                                                                               responses
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Cat IgE heavy chain amino acid sequence SEQ ID NO:25

28-MAY-2003

(first

entry)

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                                                                                                                                                                                                                    The present invention describes a method (M1) for identifying peptides CC that induce cytotoxic T-lymphocyte (CTL) response against immunoglobulin CE (IgE), comprising providing a test peptide (T) suspected of being able to bind to major histocompatibility complex (MHC) class I molecule, and cevaluating (T) for ability to elicit in a mammal a CTL response to CC induces such a response is identified. Also described are compositions: (CC) comprising at least one immunogenic peptide (I) identified by (M1); (CC) comprising at least one isolated polynucleotide encoding (I); and (CC) comprising at least one isolated polynucleotide encoding (I); and (CC) comprising antigen-presenting cells that recognise at least one (I). (CC) where CL-3 are able to bind to at least one MHC class I molecule and to CC elicit in a mammal a CTL response to naturally processed and presented CC IgE peptides. (Cl-3 can bave antiallergic, antiasthmatic, immunosuppressive, and can be used as inducers of a CTL response against IgE, and in CC vaccines. Cl-3 can be used for modulating an IgE-mediated condition in a mammal. Cl-3 can be used for modulating an IgE-mediated condition such as IgE-mediated atopic hypersensitivity conditions, IgE-mediated condition such as IgE-mediated atopic hypersensitivity conditions (Such as anaphylaxis), non-atopic catopic hypersensitivity conditions (Such as anaphylaxis), non-atopic catopic hypersensitivity conditions (Such as anaphylaxis), non-atopic catopic hypersensitivity conditions (Such as anaphylaxis), non-atopic catopic hypersensitivity conditions (Such as anaphylaxis), non-atopic hypersensitivity conditions (Such as anaphylaxis), non-atopi
                                                                                                               Query Match
Best Local
                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunoglobulin E; vaccine; IgE; cytotoxic T lymphocyte response; immune response; major histocompatibility complex; MHC; immunogenic; antiallergic; antialtergic; immunosuppressive; vasotropic; cytostatic; dermatological; antiinflammatory; IgE-mediated condition; food allergy; atopic hypersensitivity condition; allergic rhinitis; allergic asthma; atopic dermatitis; non-atopic hypersensitivity condition; anaphylaxis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying peptides that induce cytotoxic T-lymphocyte, CTL response against IgE, by identifying peptide eliciting CTL response to IgE peptides naturally presented by major histocompatibility complex class
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-2001; 2001US-0312120P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-AUG-2002; 2002WO-US026986.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; Page 145-147; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (IGET-) IGE THERAPEUTICS INC
                                                                                                                                                                                      Sequence
364
                                                                                           21;
                                                                                                                    Similarity
                                                                                                                                                                                         496 AA;
                                            CRVTHPHLPKDIVRSIAKAPGKRAP 25
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                                                                                             Conservative
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                                                                                                                 84.6%;
                                                                                                                 Score 115;
Pred. No. 6.
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                                                                                             Mismatches
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                                                                                                                                            DB 6;
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Search completed: February 22, Job time: 165 secs

2005, 06:57:14

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CKVTHPDLPKDIVRSIAKAPGRRFP

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364

RESULT 15 ABU09338

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Query Match
Best Local S
Matches
                                                                               The present invention relates to the isolation of feline immunoglobulin (IgE) kappa light chain and IgE epsilon heavy chain proteins, and the polynucleotide sequences encoding them. The sequences of the invention are useful for treating feline IgE-mediated immune responses (e.g. allergies, parasitic infections or neoplasia), in vaccine technology, small molecule/antibody technology, molecular biology, and various immunological techniques related to feline IgE and its functions. The present sequence represents feline IgE epsilon heavy chain #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reline; immunoglobulin E; IgE epsilon heavy chain; parasitic infection; IgE-mediated immune response; allergy; neoplasia; vaccine technology; antibody technology; antiallergic; antiparasitic; cytostatic.
                                                                                                                                                                                                                                             New nucleic acid molecule encoding feline immunoglobulin E (IgE) heavy light chain protein, useful for treating feline IgE-mediated responses e.g. allergies, parasitic infections or neoplasia.
                                                                                                                                                                                                                                                                                                       WPI; 2003-391997/37.
N-PSDB; ABX95715.
                                                                                                                                                                                                                                                                                                                                                                                                                           07-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JAN-2000; 2000US-00479614.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003013183-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Feline IgE
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                                                        Sequence 496
                                                                                                                                                                                                                   Claim 1; Page 37-39; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                  Mccall C,
                                                                                                                                                                                                                                                                                                                                                                               (MCCA/) MCCALL C. (WEBE/) WEBER E.
               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               epsilon heavy chain #2
 Conservative
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                84.6%;
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                Score 115; DB 6;
Pred. No. 6.2e-09;
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   Mismatches
                              Length 496
      Indels
      <u>,</u>
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Result
No.
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-Maximum Match 100%
-Tisting first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/backtTles1.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           February 22, 2005, 06:50:48 ; Search time 42 Seconds (without alignments) 44.434 Million cell updates/sec
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136
      CRVTHPHLPKDIVRSIAKAPGKRAP
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Copyright (c) 1993 - 2005 Compugen Ltd.
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   US-09-701-623C-2
US-08-336-583-2
PCT-US-95-13795-2
US-09-281-7608-39
US-09-281-7608-36
US-09-479-614-29
US-09-479-614-29
US-09-479-613C-26
US-09-701-623C-26
US-09-701-623C-27
US-09-701-623C-27
US-09-701-623C-27
US-09-701-623C-31
US-09-802-095-31
US-09-802-095-31
US-09-802-095-31
US-09-281-7608-28
US-09-281-7608-28
US-09-281-7608-24
US-09-281-7608-24
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Sequence 2, Appli
Sequence 2, Appli
Sequence 39, Appli
Sequence 36, Appli
Sequence 14, Appli
Sequence 29, Appli
Sequence 29, Appli
Sequence 26, Appli
Sequence 26, Appli
Sequence 26, Appli
Sequence 27, Appli
Sequence 31, Appli
Sequence 31, Appli
Sequence 1, Appli
Sequence 27, Appli
Sequence 28, Appli
Sequence 29, Appli
Sequence 29, Appli
Sequence 27, Appli
Sequence 4, Appli
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Query M Best Lo Matches Qy Db	LENGTH PO ORGANISM ORGANISM ORGANISM PEATURE: OTHER IN PUBLICATI P	RESULT 1 US-09-701-62: Sequence 2 Patent No. GENERAL INI APPLICANT TITLE OF TITLE OF TITLE OF FILE REFE CURRENT FI CURRENT FI PRIOR APPI	αα.αααααααααααααααααααααααααααααααααα
atch cal s 25 1	FRIDE SM: 3	623C O. 6 O. 6 O. 6 O. 6 O. 6 O. 6 O. 6 O. 6	
100.0%; Scor Similarity 100.0%; Pred 5; Conservative 0; Mi CRVTHPHLPKDIVRSIAKAPGKRAP [CRVTHPHLPKDIVRSIAKAPGKRAP CRVTHPHLPKDIVRSIAKAPGKRAP	Dog COMMATION: CH2C ORMATION: CH2C ORMATION		0.000000000000000000000000000000000000
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e 136; DB 4 . No. 2.7e-1 smatches 25 216	f dog IgE	09701623C ng Y1 B COMPOSITION AS IMMUNOGEN GY 1. US/09/701,623C 1-12-01,712-01 PCT/US99/13959 -21 09/100,287	US-09-701-623C-7 US-08-466-151-1 US-08-466-151-1 US-09-281-760E-30 US-09-100-4148-95 US-09-701-628C-92 US-09-701-628C-92 US-09-701-628C-92 US-09-100-4148-99 US-09-100-4148-99 US-09-100-4148-99 US-09-303-323-98 US-09-303-323-100 US-09-303-323-100 US-09-770-014-98 US-09-770-014-100 ALIGNMENTS
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RESULT 2
US-08-336-583-2
i Sequence 2, Application US/08336583
i Patent No. 5629415
i GENERAL INFORMATION:
APPLICANT: HOLLIS, GREGORY F.
APPLICANT: HOLLIS, MAYUR D.
ITILE OF INVENTION: DNA ENCODING CANINE IMMUNOGLOBULIN NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:

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RESULT 3
PCT-US95-13795-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application PC/TUS9513795
GENERAL INFORMATION:
APPLICANT: HOLLIS, GREGORY F.
APPLICANT: PATEL, MAYUR D.
TITLE OF INVENTION: DNA ENCODING CANINE IMMUNOGLOBULINS
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                  COUNTRY: USA
ZIP: 07065-0907
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/13795
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
TELEPHONE: (908) 594-6734
TELEPAX: (908) 594-4720
INFORMATION FOR SEQ ID NO: 2:
                                 CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: CARTY, CHRISTIME E.
REGISTRATION NUMBER: 36,099
REFERENCE/DOCKET NUMBER: 1921
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-6734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: CARTY, CHRISTINE B.
REGISTRATION NUMBER: 36,099
REFERENCE/DOCKET NUMBER: 1921
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-6734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (908) 594-4720 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/336,583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: CHRISTINE E. CARTY
STREET: 126 E. LINCOLN AVENUE; P.O. BOX 2000
CITY: RAHWAY
                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 426 amino acids
TYPE: amino acid
TYPE: amino acid
TRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 07065-0900
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126 E. LINCOLN AVENUE
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ilarity 100.0%;
Conservative 0
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Pred. No. 3.9e-13;
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APPLICANT: Mermer, Brion
APPLICANT: Prancoeur, Greg
ITILE OF INVENTION: Specific Binding Protein for Treating
ITILE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
CURRENT APPLICATION NUMBER: US/09/281,760E
CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 09/058,331
PRIOR APPLICATION NUMBER: 09/058,331
PRIOR RILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
SOPTWARE: FastSEQ for Windows Version 3.0
ISQ ID NO 39
LENGTH: 71
TYPE: PRT
ORGANISM: Canis familiaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
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                                                                                                                                           NUMBER OF SEQ ID NOS: 39
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-281-760E-39
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                                                                                                                                                                                               APPLICANT: Francoeur, Greg
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
CURRENT APPLICATION NUMBER: US/09/281,760E
CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 09/58,331
PRIOR FILING DATE: 1998-04-09
                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 36, Application US/09281760E
Patent No. 6734287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match

84.6%; Score 115; I

Best Local Similarity 100.0%; Pred. No. 9.

Matches 21; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lawton, Robert APPLICANT: Mermer, Brion APPLICANT: Francoeur, Gre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 39, Appl:
Patent No. 673428
NAME/KEY: misc_feature
LOCATION: (136)...(136)
OTHER INFORMATION: "n" stands for any nucleic acid
                                                                        TYPE: PRT
ORGANISM: Canis familiaris
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lawton, Robert
                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                     107
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NAME/KEY: misc feature LOCATION: (413)..(414) OTHER INFORMATION: "n"

stands for any nucleic acid

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APPLICANT: Weber, Eric
APPLICANT: Weber, Eric
TITLE OF INVENTION: Feline Immunoglobulin E Mol
FILE REFERENCE: P-1047
CURRENT APPLICATION NUMBER: US/09/479,614
CURRENT FILING DATE: 2000-01-07
EARLIER APPLICATION NUMBER: 60/115,033
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 14
LENGTH: 431
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; NAME/KEY: misc_feature
; LOCATION: (1832)...(1832)
; OTHER INFORMATION: "n" stands for any nucleic acid
US-09-281-760E-36
                                                                                                                                                                                                                                                                                                            US-09-479-614-14
                                US-09-479-614-14
                                                                                                                                                                                                                                                            Sequence 14, Application US/09479614
Patent No. 6573372
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 84.6%; Sc
Best Local Similarity 100.0%; P
Matches 21; Conservative 0;
 Query Match
                                             ORGANISM: Pelis catus
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OTHER INFORMATION: "n" stands
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LOCATION: (853)..(853)
OTHER_INFORMATION: "n" stands
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LOCATION: (460)..(462)
OTHER INFORMATION: "n"
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LOCATION: (451)...(451)
OTHER INFORMATION: "n"
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LOCATION: (847)..(849)
OTHER_INFORMATION: "n"
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OCATION: (1382)..(138
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; Pred. No. 1.
0; Mismatches
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 Score 115;
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ches 0;
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Length 431;
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APPLICANT: McCall, Catherine
APPLICANT: Weber, Eric
TITLE OF INVENTION: Feline Immunoglobulin E
FILE REFERENCE: P-1047
CURRENT APPLICATION NUMBER: US/09/479,614
CURRENT FILING DATE: 2000-01-07
EARLIER APPLICATION NUMBER: 60/115,033
EARLIER APPLICATION NUMBER: 34
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2
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US-09-479-614-29
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                                                                                                                                      ; TYPE: PRT; ORGANISM: Felis catus US-09-479-614-29
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                                                                Query Match 84.6
Best Local Similarity 84.6
Matches '21; Conservative
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Best Local Similarity 84.0%;
Matches 21; Conservative
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Matches 21; Conservative
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CURRENT FILING DATE: 2000-01-07
EARLIER APPLICATION NUMBER: 60/115,033
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                APPLICANT: McCall, Catherine
APPLICANT: Weber, Eric
TITLE OF INVENTION: Feline Immunoglobulin E
FILE REFERENCE: P-1047
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364 CKVTHPDLPKDIVRSIAKAPGRRFP 388
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                           1 CRVTHPHLPKDIVRSIAKAPGKRAP 25
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                                                                                  84.6%; Score 115; DB 4;
84.0%; Pred. No. 9.1e-10;
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Pred. No. 9.1e-10;
2; Mismatches 2
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2; Mismatches 2;
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RESULT 9
US-09-281-760E-26
; Sequence 26, Application US/09281760E
; Patent No. 6734287
; GENERAL INFORMATION:

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RESULT 11
US-09-701-623C-26
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PALENT NO. 6811782

FALENT NO. 6811782

GENERAL INFORMATION:
APPLICANT: Wang Ph.D., Chang Yi

TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF
TITLE OF INVENTION: ALLERGY
FILE REFERENCE: 11514153US1
CURRENT APPLICATION NUMBER: US/09/701,623C
CURRENT FILING DATE: 2000-12-01
CURRENT FILING DATE: 2000-12-01
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US-09-701-623C-6
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                                                                                                                                                                                    Sequence 26,
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Best Local (
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SEQ ID NO 6
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TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF
TITLE OF INVENTION: ALLERGY
FILE REFERENCE: 11514153US1
CURRENT APPLICATION NUMBER: US/09/701,623C
CURRENT APPLICATION NUMBER: PCT/US99/13959
PRIOR APPLICATION NUMBER: PCT/US99/13959
PRIOR FILING DATE: 1999-06-21
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR PILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
PRIOR PILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/281,760E
CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 09/058,331
PRIOR PILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
NUMBER OF SEQ ID NOS: 39
NUMBER OF SEQ ID NOS: 39
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ORGANISM: Artificial Sequence
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APPLICANT: Francoeur, Gr
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                                                                                                                                                                  6, Application US/09701623C 6811782
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100.0%; Pred. No.
tive 0; Mismatc
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100.0%; Pred. No.
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O. 3.4e-07;
O. 0;
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Sequence 27, Application US/09701623C

Patent No. 6811782

GENERAL INFORMATION:

APPLICANT: Wang Ph.D., Chang Yi

ITILE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF

PILE REFERENCE: 11514153US1

CURRENT APPLICATION NUMBER: US/09/701,623C

CURRENT FILING DATE: 1000-12-01

PAIOR APPLICATION NUMBER: DS/09/13959

PRIOR APPLICATION NUMBER: 09/100,287

PRIOR APPLICATION NUMBER: 09/100,287

PRIOR FILING DATE: 1998-06-20

NUMBER OF SEQ ID NOS: 91

SOFTWARE: Patentin Ver. 2.1
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US-09-701-623C-27
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US-09-701-623C-90
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US-09-701-623C-26
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PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 90
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Best Local (
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APPLICANT: Wang Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATE: Wang Ph.D., Chang Yi
TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF
TITLE OF INVENTION: ALLERGY
FILE REFERENCE: 11514153US1
CURRENT APPLICATION NUMBER: US/09/701,623C
CURRENT APPLICATION NUMBER: P072-01
PRIOR APPLICATION NUMBER: P072-01
PRIOR APPLICATION NUMBER: P072-01
PRIOR APPLICATION NUMBER: P072-06-21
PRIOR PORTON NUMBER: P093-06-21
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Best Local Similarity
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PRIOR APPLICATION NUMBER: 09/100,287
PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
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ORGANISM: Artificial Sequence
FEATURE:
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Pred. No.
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SEQ ID NO 27

TYPE: PRT ORGANISM: Artificial Sequence

synthetic

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OTHER IMPORMATION: Description of Artificial Sequence: sy
NAME/KEY: MOD_RES
LOCATION: (1)
OTHER IMPORMATION: I, M, L
NAME/KEY: MOD_RES
LOCATION: (2)
OTHER IMPORMATION: S, T
NAME/KEY: MOD_RES
LOCATION: (7)
LOCATION: (7
                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/100,287
PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 88
LENGTH: 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6811782
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
Chang Yi
TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF
TITLE OF INVENTION: ALLERGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 88, Application US/09701623C Patent No. 6811782
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                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/701,623C CURRENT FILING DATE: 2000-12-01 PRIOR APPLICATION NUMBER, PCT/US99/13959 PRIOR FILING DATE: 1999-06-21
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
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OTHER INFORMATION:
NAME/KEY: MOD_RES
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OTHER INFORMATION: I,
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OTHER INFORMATION: V,
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CURRENT APPLICATION NUMBER: US/09/701,623C
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: PCT/US99/13959
PRIOR FILING DATE: 1999-06-21
PRIOR PILING DATE: 1999-06-21
PRIOR PILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 87
LENGTH: 62
TYPE: PRT
ORGANISM: Artificial Sequence
PRIOR INFORMATION: Description of Artificial Sequence: synthetic
US-09-701-623C-87
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Search completed: February 22, 2005, 07:01:02 Job time : 43 secs
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Patent No. 6811782

GENERAL INFORMATION:

APPLICANT: Wang Ph.D., Chang Yi

TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF TITLE OF INVENTION: ALLERGY

FILE REFERENCE: 11514153USI
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Best Local Similarity
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                                                                       45 RVTHPHLPKDIVRSIAK 61
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100.0%; Pred. No.
                                                                                                                                                            100.0%;
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100.0%; Pred. No.
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Maximum Match 100%
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Maximum DB
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Perfect score:
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2: /cgn2_6/pcodata/2/pubpaa/PCT_NEW_PUB.pep:*

3: /cgn2_6/pcodata/2/pubpaa/US06_PUBCOMB.pep:*

4: /cgn2_6/pcodata/2/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/pcodata/2/pubpaa/US07_NEW_PUB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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/ Cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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US-09-401-636-11

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Sequence 1, Appli
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Sequence 10, Appli
Sequence 11, Appli
                                                                                                                                                                                                                                                                                                         Description
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CURRENT FILING DATE: 2001-08-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4
LENGTH: 25
TYPE: PRT
ORGANISM: DOG CH3/CH4 PEPTIDE SEQUENCE
US-09-938-700-4
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Patent No. US20020064525A1
GENERAL INFORMATION:
APPLICANT: MOZSEY, et al.
TITLE OF INVENTION: Anti-IgE Vaccines
FILE REFERENCE: PC10761A
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-10-438-7	-10-438-7	-10-453	-10-673-5	US-10-176-664-10	401-636-1	0-214-524	US-10-453-915-8	-438-794-	US-10-673-594-8	US-10-176-664-8	US-09-401-636-8	-10-214	-10-451-078-	US-10-451-078-2	-10-673-594-	US-10-673-594-4	US-10-176-664-9	US-10-176-664-4	US-09-401-636-9	US-09-401-636-4	US-10-214-524-34	US-10-152-190-14	US-10-673-594-2	-10-176-664-	US-09-401-636-2	US-10-152-190-5	-10-409-772-	US-10-409-772-2	214-524-	-09-479-614-2	-614-
	12,	equence 21,	10,	e 10,	10,	33	8	,	,	,	œ -	Sequence 30, Appl	e 4,	2	9,	4.	,	9	9	Sequence 4, Appli	u	e 1	N	e 2	,	տ	Ŋ	9	Ø	e 2	e 2

ALIGNMENTS

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RESULT 2
US-10-152-190-1
Sequence 1, Application US/10152190
publication No. US20030096369A1
GENERAL INFORMATION:
APPLICANT: Morsey, Mohamad A.
TITLE OP INVENTION: NO. US20030096369A1-anaphylactogenic IgE vaccines
FILE REFERENCE: PC11011A
CURRENT APPLICATION NUMBER: US/10/152,190
                                                                                                                                                                                                                   25;
                                                                                                                                                                                                                                                                                            n 100.0%; Score 136; DB 9; Similarity 100.0%; Pred. No. 4.6e-12; 25; Conservative 0; Mismatches n.
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Length 25;

2002-05-21

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CURRENT APPLICATION NUMBER: US/10/152,190
CURRENT FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 117
TYPE: PRT
ORGANISM: Human CH3/dog CH3 domain fusion
US-10-152-190-2
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US-10-152-190-2
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US-10-152-190-3
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US-10-152-190-3
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                                                                        Matches
                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: MOTSEY, MOhamad A.
TITLE OF INVENTION: NO. US200:
FILE REFERENCE: PC11011A
                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/10152190 Publication No. US20030096369A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 3, Application US/10152190
Publication No. US20030096369A1
GENERAL INFORMATION:
APPLICANT: Morsey, Mohamad A.
TITLE OF INVENTION: No. US20030096369A1-anaphylactogenic IgE vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2002-(
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
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CURRENT APPLICATION NUMBER: US/10/152,190
CURRENT FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 115
TYPE: PRT
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TYPE: PRT
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                        CRVTHPHLPKDIVRSIAKAPGKRAP 25
CRVTHPHLPKDIVRSIAKAPGKRAP 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CRVTHPHLPKDIVRSIAKAPGKRAP 113
                                                           100.0%; Score 136; DB 14; larity 100.0%; Pred. No. 2.5e-11; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                            US20030096369A1-anaphylactogenic IgE vaccines
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Pred. No. 2.5e-11;
                                                                                             DB 14;
                                                                                           Length 117;
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RESULT 5

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RESULT 6
US-10-152-190-12
i Sequence 12, Application US/10152190
i Publication No. US20030096369A1
i CENERAL INFORMATION:
i APPLICANT: Morsey, Mohamad A.
i TITLE OF INVENTION: NO. US2003096369A1-anaphylactogenic IgE vaccines
i FILE REFERENCE: PC11011A
i CURRENT APPLICATION UNMBER: US/10/152,190
i CURRENT FILING DATE: 2002-05-21
i NUMBER OF SEQ ID NOS: 28
i SOOTWARE: Patentin Ver. 2.1
i SEQ ID NO 12
i LENGTH: 347
i TYPE: PRT
i ORGANISM: IgE-3 fusion protein
US-10-152-190-12
                                                            ; ORGANISM: IgE-2 fusion protein
US-10-152-190-11
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US-10-152-190-11
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                                                                                                    APPLICANT: MOTSEY, Mohamad A.
TITLE OF INVENTION: No. US20030096369A1-anaphylactogenic IgE vaccines
FILE REFERENCE: PC11011A
CURRENT APPLICATION NUMBER: US/10/152,190
CURRENT FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 348
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 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                   Sequence 11, Application US/10152190 Publication No. US20030096369A1 GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 25; Conservative
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Publication No. US20030096369A1
GENERAL INFORMATION:
APPLICANT: Morsey, Mohamad A.
TITLE OF INVENTION: No. US20030096369A1-anaphylactogenic IgE vaccines
FILE REFERENCE: PC11011A
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                                                                                                TYPE: PRT
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CURRENT FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 28
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                                                                                                                                                                                                                                                                                                                                                                                                                 218 CRVTHPHLPKDIVRSIAKAPGKRAP 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       217 CRVTHPHLPKDIVRSIAKAPGKRAP 241
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 100.0%;
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Score 136; DB 14;
Pred. No. 8.5e-11;
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Pred. No. 8.4e-11;
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                 Length 348;
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RESULT 10
US-10-176-664-11
; Sequence 11, Application US/10176664
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PRIOR FILING DATE: 2001-08-13
NUMBER OF SEQ ID NOS: 61
SOFTWARE: PatentIn version 3.1
SEQ ID NO 28
LENGTH: 426
                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
SOFTWARE: FASTSEQ for
SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11, Application US/09401636 Patent No. US20010038843A1
                                                                                                                                                        Query Match
Best Local Similarity
Matches 24; Conser
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TITLE OF INVENTION: ENHANCED VACCINES
FILE REFERENCE: 10223/006601
CURRENT APPLICATION NUMBER: US/09/401,636
CURRENT FILING DATE: 1999-09-22
PRIOR APPLICATION NUMBER: US 60/106,652
PRIOR APPLICATION NUMBER: US 60/106,652
PRIOR FILING DATE: 1998-11-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: IMMUNOGLOBULIN B VACCINES AND METHODS OF USE THEREOF FILE REFERENCE: IGS-00101.P.1.1 CURRENT APPLICATION NUMBER: US/10/214,524 CURRENT FILING DATE: 2002-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chen, Swey-Shen Alex APPLICANT: Yang, Yong-Min
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Dog
                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                   FEATURE: OTHER INFORMATION: Synthetically generated proteins
                                                                                                                                                                                                                                                                                                             ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   294 CRVTHPHLPKDIVRSIAKAPGKRAP 318
                                                                                         211 CRVTHPHLPKDIVRSIAKLPGKRLAP 236
                                                                                                                                                                                                                                                                                                             341
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                                                                                                                                                                        88.6%;
92.3%;
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                                                                                                                                                        Score 120.5; DB 9
Pred. No. 1.3e-08;
0; Mismatches 1
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                                                                                                                                                                                                                                                                          Best Local
Matches :
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LENGTH: 341
                                                                                                                                                                                                                                                                                                         Query Match
       URRENT APPLICATION NUMBER: US/09/479,614
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RESULT 12
US-09-479-614-14
J Sequence 14, Application US/09479614
Publication No. US20030013183A1
GENERAL INFORMATION:
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US-10-673-594-11
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US-10-176-664-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: ENHANCED VACCINES
FILE REFERENCE: 10223/006001
CURRENT APPLICATION NUMBER: US/10/673,594
CURRENT FILING DATE: 2003-09-29
PRIOR APPLICATION NUMBER: US/09/401,636
PRIOR FILING DATE: 1999-09-22
PRIOR APPLICATION NUMBER: US 60/106,652
PRIOR FILING DATE: 1999-11-02
PRIOR FILING DATE: 1998-11-02
PRIOR FILING DATE: 1998-11-02
PRIOR FILING DATE: 1998-11-02
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CURRENT APPLICATION NUMBER: US/10/176,664

CURRENT FILING DATE: 2002-06-19

PRIOR APPLICATION NUMBER: US/09/401,636

PRIOR FILING DATE: 1999-09-22

PRIOR APPLICATION NUMBER: US 60/106,652

PRIOR FILING DATE: 1998-11-02

PRIOR FILING DATE: 1998-11-02

PRIOR FILING DATE: 1988-11-02

PRIOR FILING DATE: 1988-11-02

PRIOR FILING DATE: 1988-11-02

NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11, Application US/10673594 Publication No. US20040076625A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                                       APPLICANT: McCall, Cath
APPLICANT: Weber, Eric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hellman, Lars T.
TITLE OF INVENTION: ENHANCED VACCINES
ITTLE OF INVENTION: Feline Immunoglobulin E Molecules and Related Methods
FILE REFERENCE: P-1047
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Hellman, Lars T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                   211 CRVIHPHLPKDIVRSIAKLPGKRLAP 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      211 CRVTHPHLPKDIVRSIAKLPGKRLAP 236
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                                                                                                                                                                                                                                                                           1 CRVTHPHLPKDIVRSIAKAPGKR-AP 25
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                                                                 Catherine
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Pred. No. 1.3e-08;
0; Mismatches 1
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Pred. No. 1.3e-08;
0; Mismatches 1
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CURRENT APPLICATION NUMBER: US/09/479,614
CURRENT FILING DATE: 2000-01-07
EARLIER APPLICATION NUMBER: 60/115,033
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEO ID NOS: 34
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2
SEQ ID NO 2
LENGTH: 496
TYPE: PAT
CORGANISM: Felis catus
US-09-479-614-2
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EARLIER APPLICATION NUMBER: 60/115,033

EARLIER FILING DATE: 1999-01-07

NUMBER OF SEQ ID NOS: 34

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 14

LENGTH: 431

TYPE: PRT
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US-10-409-772-14
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US-09-479-614-14
                                                                                                                                                                                                                                                           Sequence 2, Application US/09479614
Publication No. US20030013183A1
GENERAL INFORMATION:
APPLICANT: McCall, Catherine
APPLICANT: Weber, Eric
TITLE OF INVENTION: Feline Immunoglobulin E Molecules and Related Methods
FILE REFERENCE: P-1047
   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/409,772
CURRENT FILING DATE: 2003-04-07
PRIOR APPLICATION NUMBER: US/09/479,614
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 14
TYPE: DATE: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/10409772
Publication No. US20030216565A1
GENERAL INFORMATION:
APPLICANT: McCall, Catherine
APPLICANT: WcCall, Catherine
APPLICANT: WcBer, Bric
TITLE OF INVENTION: Feline Immunoglobulin E Molecules and Related Methods
FILE REFERENCE: P-1047
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Best Local Similarity
Matches 21; Conserv
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Best Local
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Local Similarity 84.0%;
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   84.6%;
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Pred. No. 9.7e-08;
2; Mismatches 2;
Score 115; DB 10;
Pred. No. 1.1e-07;
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Pred. No. 9.7e-08;
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                 Length 496;
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CURRENT APPLICATION NUMBER: US/09/479,614
CURRENT FILING DATE: 2000-01-07
EARLIER APPLICATION NUMBER: 60/115,033
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 29
LENGTH: 496
TYPE: PAT
ORGANISM: Felis catus
US-09-479-614-29
Search completed: February 22, 2005, 07:14:10 Job time : 132 secs
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US-09-479-614-29
; Sequence 29, Application US/09479614
; Publication No. US20030013183A1
; GENERAL INFORMATION:
; APPLICANT: McCall, Catherine
; APPLICANT: McCall, Catherine
; TITLE OF INVENTION: Feline Immunoglobulin E Molecules and Related Methods
; FILE REFERENCE: P-1047
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                                                                                                                            Query Match 84.6%;
Best Local Similarity 84.0%;
Matches 21; Conservative
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Pred. No. 1.1e-07;
2; Mismatches 2;
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Database: PIR 79;* 1: pir1:* 2: pir2:* 3: pir3:* 4: pir4:*	Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries	Minimum DB seq length: 0 Maximum DB seq length: 2000000000	Total number of hits satisfying chosen parameters: 283416	Searched: 283416 seqs, 96216763 residues	Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5	Title: US-10-751-743-4 Perfect score: 136 Sequence: 1 CRVTHPHLPKDIVRSIAKAPGKRAP 25	Run on: February 22, 2005, 06:51:23 ; Search time 38 Seconds (without alignments) 63.301 Million cell updates/sec	OM protein - protein search, using sw model	GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
			283416			•	<pre>38 Seconds nments) n cell updates/sec</pre>		

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	Ø	ຫ	4	ω	N	۲	No.	Result
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hypothetical prote	Ig gamma-2a chain	Ig gamma-2a chain	Ig gamma-2a chain	Ig gamma-2a chain	Ig gamma 2a chain	ain	Ig gamma 3 chain c	hai	ain	a 4 cl	myosin heavy chain	n heavy o	cardiac myc	Ig Y heavy chain (prot	hypothetical prote	myosin heavy chain	myosin alpha heavy	ᅜ	Ig heavy chain pre	novel antigen rece	B chain C3 regi	B chain C3 regi	epsilon ch	epsilon chain	epsilon chain	-chain	epsilon	Description	

45	44	43	42	41	40	39	38	. 37	36	35	34	33	32	31	30
48.5	48.5	48.5	48.5	48.5	48.5	49	49	49.5	49.5	49.5	50	50	50	50	50
35.7	35.7	35.7	35.7	35.7	35.7	36.0	36.0	36.4	36.4	36.4	36.8	36.8	36.8	36.8	36.8
374	330	329	255	234	166	1938	1937	573	474	405	1935	1935	1935	1934	1038
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869339	GHHU	800847	831866	PT0207	C72641	JX0178	I38055	612838	G2MS11	G2MSBM	A59286	806006	A37102	148153	MWRBCB
Ig heavy chain V r	Ig gamma-1 chain C	Ig gamma-2c chain	Ig gamma-1 chain C	Ig gamma chain C r	hypothetical prote	myosin heavy chain	myosin heavy chain	Ig mu chain precur	Ig gamma-2b chain	Ig gamma-2b chain	myosin heavy chain	myosin beta heavy	myosin beta heavy	myosin heavy chain	myosin beta heavy

ALIGNMENTS

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A; Molecule type: mRNA
A; Residues: 'N', 169-307, 'L', 309-342 < KIN>
A; Residues: 'N', 169-307, 'L', 309-342 < KIN>
C; Complex: An immunoglobulin heterotetramer subunit consists of two identical light (ks. c) complex: An immunoglobulin hemotes, such as IgA and IgM, the subunits associate into l. c; Superfamily: immunoglobulin C region; immunoglobulin homology
C; Superfamily: immunoglobulin homology < IM1>
F; 19-80/Domain: immunoglobulin homology < IM1>
F; 118-186/Domain: immunoglobulin homology < IM2>
F; 223-291/Domain: immunoglobulin homology < IM3>
F; 327-398/Domain: immunoglobulin homology < IM3>
F; 346,99,170,240,265,369,419/Binding site: carbohydrate (Asn) (covalent) #status predict.
C;Accession: 136948
R;Sakoyama, Y.; Hong, K.
Proc. Natl. Acad. Sci. U.S.A. 84, 1080-1084, 1987
Proc. Natl. Acad. Sci. U.S.A. 84, 1080-1084, 1987
A;Title: Nucleotide sequences of immunoglobulin epsilon genes of chimpanzee and oran A;Reference number: 136948; MUID:87147196; PMID:3103123
                                                                                                                                                                                                                                                                                                                                    RESULT 2
I36948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig epsilon chain C region - rat (Species: Rattus norvegicus (Norway rat))
(Species: Rattus norvegicus (Norway rat))
(Species: T-Dec-1982 #sequence revision 17-Dec-1982 #text_change 09-Jul-2004
(C;Date: 17-Dec-1982 #sequence revision 17-Dec-1982 #text_change 09-Jul-2004
(C;Accession: A93442; A90937; Ā02143
(C;Accession: A93442; A90937; A.; Karlsson, T.; Bennich, H.

Nucleic Acids Res. 10, 6041-6049, 1982
A;Title: Structure and evolution of the heavy chain from rat immunoglobulin E.
A;Reference number: A93442; MUID:83064537; PMID:6292865
                                                                                                                                                                                                Ig epsilon-chain - chimpanzee (fragment)
C;Species: Pan troglodytes (chimpanzee)
C;Date: 04-Oct-1996 #sequence_revision 04-Oct-1996 #text_change 21-Jan-2000
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A;Contentes myeloma IR162
A;Accession: A90937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:P01855
A;Experimental source: strain LOU/c/Wsl, immunocytoma IR2
R;Kindsvogel, W.R.; Reddy, E.P.; Moore, J.M.; Faust Jr., (
DNA 1, 335-343, 1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-429 < HEL>
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Best Local Similarity
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Pred. No. 2.6e-07;
1; Mismatches 4; Indels
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A.Rosiques type: DNA
A.Rosiques 1-358,'L',360-428 <MAX>
A.Cross-references GB.U00222, NID:9184755
A.Roste: this sequence difference may be due to polymorphism
A.Crossin: MH.J. Johansson, S.G.O., Von Bahr-Lindstrom, H.
A.Acceston: Ab4418 A94418
A.Rosterin: A944
                                                                                                                                                                                                                                                                                                                                                                                                                     A)Cross-reference CRANA MID:9184755

A)Cross-reference GB:000222, NID:9184755

A)Fittle: Two unusual forms 98 E.

A)Reference number: PHI214; MUID:92308839; PMID:1613458

A)Cross-references EMBL:N6393; GB:838668; NID:932987

A)Cross-references EMBL:N6393; GB:838668; NID:932987

A)Fittle: Noiseular forms 92 GB:838668; NID:932987

A)Fittle: Noiseular forms 92 GB:838668; NID:932987

A)Fittle: Noiseular forms 92 GB:838668; NID:932987

A)ACCESSION: A93491; MUID:83168897; PMID:6300763

A)Molecule type: mRNA

A)Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                          iResidues: 1-358,
                                                                                                                                                                                                                                                           Molecule type: D
Residues: 2-428
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Best Local Similarity 66.7%;
16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Accession: 136948
A;Molatus: preliminary;
A;Rosidues: type: DNA
A;Rosidues: 1-426 <RES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Grose-references: GB:M15398; NID:g176797; PIDN:AAA35416.1; PID:g176798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ng terminal ----
| number: hal repeat-like elements flank a human immunoglobulin epsilon pseudog
| north A23195; MUID:04207910; PMID:6327276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           297 CRVTHEHLERALVRSTTKTSGERÅ 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CRVTHPHLPKDIVRSIAKAPGKRA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .: UNIPROT:P01854; GB:L00022; GB:J000227; GB:V00555; NID:9185035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              65.4%; Score 89; DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nes 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 426;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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Alcrose references: GDB:119335; OMIM:147180

Almap postion: 14922,33-14932,33

Almap postion: 14922,33-14932,33

Lintrons: 11: 104/1; -311/1; 319/1

C.Complex: Al: 104/1; -311/1; 319/1

C.Superfamily: immunoglobulin heterotetramer subunit consists of two identical light (Reywords: alternative policing, dupn; immunoglobulin homology and IgM, the subunits associate into 1 F:22-301/Domain: immunoglobulin homology <IM1; 919/10001: immunoglobulin homology <IM2; F:338-407/Domain: immunoglobulin homology <IM2; F:138-407/Domain: immunoglobulin homology <IM2; F:15-105,20 bonds: interchain homology <IM2; F:121,49.99-184,513-193,239-293in (to 119th chain)

F:121,299/146(252,275/Binding site: 03/Disulfide bonds: #Status predicted

F:121,209/Disulfide bonds: interchain (to heavy chain) #Status predicted

experimental
Query Match

Best Local Similarity 63.2%; Score 86; DB 1; Length 428;

Matches 15; Conservative 3; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                      A;Molecule (170e; DNA)
A;Residues; 401-428 <4B3
A;Experimented B:853497; NID:9263162; PIDN:AAB24855.1; PID:9263163
A;Note: Bequence extracted from NCBI backbone (NCBIP:123483)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Status: Preliminary, not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Notecule type: mRNA,

Notecule type: mRNA,

Notecule type: mRNA,

Notecule type: mRNA,

Cross-references: GB:655276, NID:9263168, PIDN:AAB24858.1, PID:9263169, NOtec: Bequence: B cell myeloma U-266

NOtec: Bequence extracted from NCBI backbone (NCBIP:125299)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Cross references; GB:855273; NID:9263166; PIDN:AAB24857.1; PID:9263167; Accession; D46536 from NCBI backbone (NCBIP:125297)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hellman, L.

VILLE: Characterization of four novel epsilon chain mRWA and a comparative analogues. PMID: 93122085; PMID: 8419166

A.Accession: C46536; MUID: 93122085; PMID: 8419166

A.Accession: Traitminary; not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           J. Biol. Chem. 269 8.; Cheah, H.K.; Saxon, A.
A./Title: Complex alternative RNA splicing of epsilon-immunoglobulin transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'Molecule type: mRy
'Residues: 98-352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Kenten, J.H.; Molgaard, H.V.; Houghton, M.; Derbyshire, R.B.; Viney, J., Beil, L.O., A;Ritle: Cloning and Bequence determination of the gene for the human immunoglobulin a.a.roesion. Rozozz A9393; MUID:83065234; PMID:6815656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A, Molecule type; mRNA
A, Residues; 1-40;68-114,427-428 <KEN,
A, Cross references; GB;L00022; NID;9185035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1411e: Purification and characterization of a recombinant human IgE Fc-eps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Preliminary, not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ule type: mRNA
(uss: 320-428 <ZH2>
'Amental source: myeloma U266-derived cell line AP-10
nan. L. Chiracted from NCBI backbone (NCBIN:141701, NCBIP:141702)
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RESULT 6
168726
198 Chain C3 region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Aug-1996 #sequence_revision 02
C;Accession: 168726
R;Shinkai, Y; Makauchi, H.; Honjo, T.; O
Immunogenetics 27, 288-292, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:Z27397; NID:g416537; PIDN:CAA81788.1; PID:g940782
C;Superfamily: immunoglobulin C region; immunoglobulin homology
F;353-421/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Description: Combination A;Reference number: $38864 A;Accession: $38864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Kipp, B.; Becker, W.; Schlaak, M. submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ig epsilon chain C region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 24-May-2001
C;Accession: 638864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPROT: P06336; GB:J00476; NID:g194875; PIDN:AAA38085.1; PID:g387220; C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kar hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin C region; immunoglobulin homology C;Keywords: duplication; glycoprotein; heterotetramer; immunoglobulin C;Keywords: duplication; glycoprotein; heterotetramer; immunoglobulin F;1-44/Domain: immunoglobulin homology (fragment) <IM1>
F;81-149/Domain: immunoglobulin homology <IM2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Accession: A02144
R;Liu, F.T.; Albrandt, K.; Sutcliffe, J.G.; Katz, D.H.
Proc. Natl. Acad. Sci. U.S.A. 79, 7852-7856, 1982
A;Title: Cloning and nucleotide sequence of mouse immunoglobulin epsilon chain A;Reference number: A02144; MUID:83117774; PMID:6818553
A;Accession: A02144
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A; Residues: 1-548 <KIP>
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F;290-361/Domain: immunoglobulin homology <IM4>
F;10,51,62,133,205,228,332,382/Binding site: carbohydrate (Asn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig epsilon chain C region (version 1) - mouse (fragment)
C/Species: Mus musculus (house mouse)
C/Date: 13-Jun-1983 #sequence_revision 13-Jun-1983 #text_change 09-Jul-2004
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A;Residues: 1-388 <LIU>
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Best Local
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Best Local Similarity
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; Pred. No. 0.00
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                           Score 80; DB 2;
Pred. No. 0.0013;
2; Mismatches
                             T., Okumura,
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                                                                             02-Aug-1996 #text_change
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0.00094;
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RESULT S04845

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novel antigen receptor precursor - nurse shark (,Species: Ginglymostoma cirratum (nurse shark) (,Species: Ginglymostoma cirratum (nurse shark) (,C;Date: 10-Apr-1996 #sequence_revision 19-Apr-1996 #text_change 09-Jul-2004 (,Accession: S60266 R;Greenberg, A.S.; Avila, D.; Hughes, M.; Hughes, A.; McKinney, B.C.; Flajnik, Nature 374, 168-173, 1995 Nature 374, 168-173, 1995 A;Reference number: S60266; MUID:95183140; PMID:7877689 A;Accession: S60266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:M22930; NID:g194455; PIDN:AAA37911.1; PID:g194460 C;Superfamily: immunoglobulin C region; immunoglobulin homology F;22-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Title: Mouse immunoglobulin allotypes: Multiple differences between the nucleic acid A;Reference number: 154443; MUID:88152907; PMID:3346043
A;Accession: I68726
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-107 <RESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Shinkai, Y.; Nakauchi, H.; Honjo, T.; Okumura, K. Immunogenetics 27, 288-292, 1988
A;Title: Mouse immunoglobulin allotypes: Multiple differences A;Reference number: I54443; MUID:88152907; PMID:3346043
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A; Residues: 1-107 < RES>
                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-684 < GRE>
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C;Superfamily: immunoglobulin C region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IgE chain C3 region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Aug-1966 #sequence_revision 02-Aug-1996 #text_change 21-Jan-2000
C;Accession: I68730
                                                                                                                                                                               A;Cross-references: UNIPROT:Q90544; EMBL:U18701; NID:g699442; PIDN:AAB48195.1; PID:g699
                                                                                                                                                                                                                                             A; Status: preliminary; nucleic acid sequence not shown
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                                                                                       Matches
                                                                                                                                  Query Match
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                                          1 CRVTHPHLPKDIVRSIAKAPGK 22
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                                                                                                             h 43.4%;
Similarity 50.0%;
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CLVGHPSLNRDLIRSTNKSNGK 666
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immunoglobulin homology
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ilarity 65.0%;
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                                                                                       4; Mismatches
                                                                                                             Score 59; DB 2; Pred. No. 1.7;
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Pred. No. 0.019;
0; Mismatches
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Pred. No. 0.019;
0; Mismatches
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A.M.; Chambers, A.P.; Nadal-Ginard,

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A;Molecule type: mRNA
A;Residues: 1512-1574,'S',1576-1721,'N',1723-1851,'N',1853-1869,'N',1871-1933,'I',1935-I
A;Note: there are 10 or more myosin heavy chain genes in the rat, at least two of which
                                                                                          A.Wolecule type: protein
A.Wolecule type: protein
R.Residues: 1-12, 'AP',14-45,'A',47-50,'AP',53-81,'E',83-86,'Q',88-109,111-133,'H',135-1(
R.Mahdavi, V.; Periasamy, M.; Nadal-Ginard, B.
Nature 297, 659-664, 1982
A.Title: Molecular Characterization of two myosin heavy chain genes expressed in the ad-
A.Reference number: A02988; MUID:82220036; PMID:7045682
A.Accession: A02988
                                                                                                                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-1938 <MC2>
R;Mahdavi, V.; Chambers, A.P.; Nadal-Ginard, B.
Proc. Natl. Acad. Sci. U.S.A. 81, 2626-2630, 1984
A;Title: Cardiac alpha- and beta-myosin heavy chain genes are organized in A;Accession: A20971; MUID:84194059; PMID:6585819
A;Accession: A20971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 1-1938 <MCN>
A;Residues: 1-1938 <MCN>
A;Cross-references: UNIPROT: P02563; EMBL::X15938; NID:956654; PIDN:CAA34064.1; R;MCNally, E.M.; Kraft, R.; Bravo-Zehnder, M.; Taylor, D.A.; Leinwand, L.A.
J. Mol. Biol. 210, 665-671, 1989
A;Title: Full-length rat alpha and beta cardiac myosin heavy chain sequences.
A;Reference number: S07535; MUID:90133919; PMID:2614840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Complete nucleotide sequence of full length cDNA for rat alpha cardiac A;Reference number: S06005, MUID:90016822; PMID:2798111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Rattus norvegicus (Norway rat)
C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: S06005; S07535; A20971; A02988; I53305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  myosin alpha heavy chain, carciac muscle [similarity] N;Alternate names: alpha-myosin heavy chain N;Contains: myosin ATPase (EC 3.6.4.1)
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A;Molecule type: mRNA
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Nucleic Acids Res. 17, 7527-7528, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Reference number: $05695
A;Accession: $05695
A;Molecule type: mRNA
A;Residues: 'LC', 3-308,'H',310-549 <LIT>
A;Cross-references: EMBL:XI5114; NID:g64799; PID:g763031
C;Superfamily: immunoglobulin C region; immunoglobulin homology
C;Keywords: glycoprotein; heterotetramer; immunoglobulin
C;Keywords: glycoprotein; heterotetramer; immunoglobulin
F;26-109/Domain: immunoglobulin homology <IMMy
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C;Species: Xenopus laevis (African clawed frog)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 26-Aug-1999
C;Accession: S04845; S05695
R;Amemiya, C.T.; Halre, R.N.; Litman, G.W.
Nucleic Acids Res. 17, 5388, 1989
A;Title: Nucleotide sequence of a cDNA encoding a third distinct Xenopus im A;Reference number: S04845; MUID:89345103; PMID:2503814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the EMBL Data Library, April 1989
A;Reference number: 805695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: mRNA
A;Residues: 1-549 <AME>
A;Cross-references: EMBL:X15:114
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12; Conserv
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52.2%;
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Pred. No. 1
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1.9;
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A;Status: preiiman,
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-32 <BRA>
A;Cross-references: EMBL:X56181; NID:g28318; PIDN:CAA39642.1; PID:g28319
A;Cross-references: EMBL:X56181; PID:g28318; PIDN:CAA39642.1; PID:g28319
A;Cross-references: EMBL:X56181; PID:g28318; PIDN:G28318; PIDN:G28318; PIDN:G28318; PIDN:G28318; PIDN:G28318; PIDN:G28318; PID:g28318;
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R;Brand, N.J.; Dabhade, N.; Yacoub, M.; Barton, P.J.R.
Biochem. Biophys. Res. Commun. 179, 1255-1258, 1991
A;Title: Determination of the 5' exon structure of the human cardiac alpha-myosin heavy
A;Reference number: S18830; MUID:92028859; PMID:1930170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: mNNA
A;Residues: 1-1461 «MA2>
R;Bpp, T. A.; Dixon, I.M.C.; Wang, H.Y.; Sole, M.J.; Liew, C.C.
R;Bpp, T. A.; Dixon, I.M.C.; Wang, H.Y.; Sole, M.J.; Liew, C.C.
A;Title: Structural organization of the human cardiac alpha-myosin heavy chain gene (MYHA;Accession: A49354; MUID:94140346; PMID:8307559
A;Accession: A49354
A;Status: nucleic acid sequence not shown; translation not shown
A;Residues: 1-87,'Q',89-573,'Q',575-607,'A',609-743,'T',745-789,'M',791-1013,'V',1015-10
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C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: A46762; B46762; Ā49354; S18830; B32562; B33835; B27858; A28908
R;Matsuoka, R; Beisel, K.W.; Furutani, M.; Arai, S.; Takao, A.
Am. J. Med. Genet. 41, 537-547, 1991
A;Title: Complete sequence of human cardiac alpha-myosin heavy chain gene and A;Accession: A46762; MUID:92133665; PMID:1776652
A;Reference number: A46762; MUID:92133665; PMID:1776652
A;Status: nucleic acid sequence not shown
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A46762
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A; Molecule type; mRNA
A; Residues: 1872-1933, 'I', 1935-1938 <RES>
A; Cross-references: GB:M32697; NID:g205596; PIDN:AAA41658.1;
A; Cross-references: GB:M32697; chain; myosin motor domain homolo
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Bur. Heart J. 5, 181-191, 1984
A;Title: Cardiac myosin heavy
A;Reference number: 153305, MU
A;Accession: 153305
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Contains: myosin ATPase (EC 3.6.4.1)
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A;Molecule type: DNA
A;Residues: 1-3,'S',5-10,'T',12,14-67 <SAB>
R;Karabayashi, M.; Tsuchimochi, H.; Komuro, I.; Takaku, F
J. Clin. Invest. 82, 524-531, 1988
A;Title: Molecular cloning and characterization of human
R;Liew, CC.; Jandreski, M.A.
R;Liew, CC.; Jandreski, M.A.
Proc. Natl. Acad. Sci. U.S.A. 83, 3175-3179, 1986
A;Title: Construction and characterization of the alpha-form
A;Reference number: A23938; MUID:86205859; PMID:3458174
                                                                                                                                                                                                                                                                                                                                     R;Wang, R.; Sole, M.J.; Cukerman, B.; Liew, C.C. J. Mol. Cell. Cardiol. 26, 1155-1165, 1994
A;Title: Characterization and nucleotide sequence of the A;Reference number: 148153; MUID:95115033; PMID:7815459
A;Accession: 148175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               myosin heavy chain alpha, cardiac muscle [similarity] - C;Species: Mesocricetus auratus (golden hamster) C;Date: 02-dul-1996 #sequence revision 02-dul-1996 #text C;Accession: 148175; A23938
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148175
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F;129/Modified site: N6,N6-trimethyllysine (Lys)
F;184/Binding site: ATP (Lys) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Superfamily: myosin heavy chain; myosin motor domain C;Keywords: actin binding; ATP; cardiac muscle; coiled F;88-768/Domain: myosin motor domain homology <a href="https://www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin.motor.com/air-nomology-www.myosin
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A;Residues: 1407-1532,'N',1534-1539,'M',1541-1576,'NV',1579-1704,'EQ',1707-1762,'D',1764
A;Cross-references: GB.M21664; NII:g189006; PIDN:AAA36344.1; PID:g386972
A;Experimental source: fetal heart
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A;Molecule type: DNA
A;Residues: 1-87,'Q',89-177;1551-1732,'E',1734-1736,'T',1738-1762,'D',1764-1939
R;Residues: LJ, Gianola, K.M.; McNally, B.M.; Peghali, R.; Eddy, R.; Shows, T.B.;
Rucleic Acids Res. 15, 5443-5459, 1987
A;Title: Human cardiac myosin heavy chain genes and their linkage in the genome.
A;Reference number: A93669; MUID:87260010; PMID:3037493
A;Recession: B27858
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A; Accession: A28908
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A; Accession: B33835
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Molecule type: DNA

;Residues: 1-87, (27, 89-177;1551-1732, E',1734-1736, T',1738-1762, D',1764-1848 <YAl>

;Residues: 1-87, (27, 89-177;1551-1732, E', 1734-1736, T',1738-1762, D',1764-1848 <YAl>

;Yamauchi-Takihara, K.; Sole, M.J.; Liew, J.; Ing, D.; Liew, C.C.

;Yamauchi-Takihara, K.; Sole, M.J.; Liew, J.; Ing, D.; Liew, C.C.
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                                                                                                                                                                       ,Molecule type: DNA
,Residues: 1-1939 <RES>
,Cross-references: UNIPROT:P13539; GB:L15351, NID:g402373, PIDN:AAB59701.1;
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Best Local (
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VTHPHLPKDIVRSIAKAPGKR

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C;Accession: B87319

R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, B; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Koln, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001

A;Title: Complete Genome Sequence of Caulobacter crescentus.

A;Reference number: A87249; MUID:21173698; PMID:11259647
                                                                                                             A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-197 <SEE> A;Cross-references: UNIPROT:Q9ZBJ8; EMBL:AL035161; A;Experimental source: strain A3(2)
                                                                                                                                                                                                                                                                                                                                                                 RESULT 14
T35963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein SC9C7.12 - Streptomyces coelicolor
C;Species: Streptomyces coelicolor
C;Date: 03-Dec_1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Introns: 67/3; 115/3; 168/1; 177/2; 214/3; 245/3; 267/1; 300/1; 33 992/2; 1453/3; 1509/1; 1550/3; 1653/3; 1751/3; 1763/3; 1855/3; 1887/ C;Supérfamily: myosin heavy chain; myosin motor domain homology C;Keywords: ATP; cardiac muscle; heart; nucleotide binding; P-loop F;88-768/Domain: myosin motor domain homology cMMOT>
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A;Molecule type: mRNA
A;Residues: 1630-1843,'R',1845-1878,'T',1880-1927,'N',1929-1932,1934-1939 <LIE>
C;Genetics:
                                                                                                                                                                                                              A; Reference number: Z21551
A; Accession: T35963
                                                                                                                                                                                                                                                    R;Seeger, K.J.; Harris, D.; Bentley, S.D.; Parkhill, submitted to the EMBL Data Library, January 1999
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A;Molecule type: DNA
A;Residues: 1-273 <STO>
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                                       Query Match
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   Similarity
9; Conserv
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 37.5%;
ilarity 42.9%;
Conservative
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Score 51; DB Pred. No. 7.1; 6; Mismatches
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Pred. No.
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Pred. No. 26;
                                                                                                                                    EMBL:AL035161; PIDN:CAA22724.1;
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                                         Length 197;
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RESULT 15

B46529

Ig Y heavy chain (7.88) - duck
NALternate names: Ig gamma chain (7.88)
C;Species: Anas platyrhynchos (domestic duck)
C;Pate: 18-Jun-1993 #sequence revision 18-Nov-1994 #text_change 21-Jul-2000
C;Accession: B46529; S20759 -
C;Magor, K.E.; Warr, G.W.; Middleton, D.; Wilson, M.R.; Higgins, D.A.
J. Immunol. 149, 2627-2633, 1992
A;Title: Structural relationship between the two IgY of the duck, Anas platyrhynchos: md
A;Reference number: A46529; MUID:93017865; PMID:1401901
A;Accession: B46529
A;Status: preliminary
A;Rolecule type: mRNA
A;Residues: 1-572 <AMGS-
A;Cross-references: EMBL:X65219; NID:962442; PIDN:CAA46322.1; PID:962443
A;Reperimental source: spleen
A;Note: sequence extracted from NCBI backbone (NCBIP:116127)
C;Superfamily: immunoglobulin C region; immunoglobulin homology
F;37-120/Domain: immunoglobulin homology <IMM>
Search completed: February 22, 2005, 07:01:47 Job time : 39 secs
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his SWISS-PROT entry is copyright. The European Bioinformatics Institute of Bio But Depart Bioinformatics Institute Be by non-profit institutions Be by non-profit institutions Be by non-profit institutions Be dodified and this statement is not Intitles requires a license agreeme Intitles requires a license disb-sil Fend an email to licensedisb-sil Fend an email to lic	[3] SEQUENCE OF 205-306 FROM N.A. MEDLINE=82174576; PubMed=6803238; Hellman L., Pettersson U., Bennich "Characterization and molecular cli (epsilon) chain of rat immunoglobu Proc. Natl. Acad. Sci. U.S.A. 79:1	Nucleic Acids Res. 10:6041-6([2] SEQUENCE OF 168-342 FROM N.A. MEDLINE=83182019; PubMed=6820 Kindsvogel W.R., Reddy E.P., "A cloned cDNA probe for rat construction, identification, DNA 1:335-343(1982).		AC P01855; DT 21-UUL-1986 (Rel. 01, Created) DT 21-UUL-1986 (Rel. 01, Last sequence update DT 10-OCT-2003 (Rel. 42, Last annotation upda DE Ig epsilon chain C region. OS Rattus norvegicus (Rat). OC Eukaryota; Metazoa; Chordata; Craniata; Ve OC Mammalia; Eutheria; Rodentia; Sciurognathi OX NCBI_TaxID=10116;	၂၀ ဗ	32 50 36.8 1934 1 MYHT MESAU 33 50 36.8 1935 1 MYHT HUMAN 34 50 36.8 1935 1 MYHT PIG 35 50 36.8 1935 2 QMYHT RAT 36 50 36.8 1935 2 QMHT RAT 37 50 36.8 1935 2 QMHT RAT 38 50 36.8 1935 2 QMHT RAT 38 50 36.8 1935 2 QMHT RAT 39 50 36.4 374 2 QMHT RAT 49 5 36.4 374 374 SPOP HUMAN 49 5 36.4 374 1 SPOP HUMAN 49 5 36.4 374 1 SPOP MOUSE 44 49.5 36.4 374 1 SPOP MOUSE 45 49.5 36.4 374 2 QMHT RAT 46 49.5 36.4 374 2 QMHT RAT 47 49.5 36.4 374 2 QMHT RAT 48 49.5 36.4 374 2 QMHT RAT 48 49.5 36.4 374 2 QMHT RAT 49 5 36.4 374
ght. It is produced through a collaboration Bioinformatics and the EMBL outstation - tritute. There are no restrictions on its no as long as its content is in no way not removed. Usage by and for commercial eement (See http://www.isb-sib.ch/announce/-sib.ch).	the mRNA f (1982).	IR162). ., Faust C.H. J bulin epsilon h sequence.";	A., Karlsson T., Bennich H., chain from rat immunoglobulin	update) nn update) sta; Vertebrata; Euteleostomi; gnathi; Muridae; Murinae; Rattus.	ENTS	U P13540 mesocricetu N P12883 homo sapien P79233 sus scrofa P02564 rattus norv O8mjus equus cabal O9be39 bos taurus O9gkr1 sus scrofa O91z83 mus musculu P01866 mus musculu P01866 mus musculu O761v9 mus musculu O6zws8 mus musculu O6p8b3 xenopus tro O7zx06 xenopus lae

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Query Match
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        SEQUENCE OF 1-40; 68-114 AND 427-428 FR MEDLINE-83065234; PubMed=6815656; Kenten J.H., Molgaard H.V., Houghton M. Bell L.O., Gould H.J.; "Cloning and sequence determination of immunoglobulin epsilon chain expressed
                                                                                                                                                 SEQUENCE FROM N.A.

MEDILINE-84207910; PubMed=6327276;
Ueda S., Nakai S., Nishida Y., Hisajima H.,
"Long terminal repeat-like elements flank a
epsilon pseudogene that lacks introns.";
EMBO J. 1:1539-1544(1982).
                                                                            Bennich H.H., Johansson S.G.O., von Bahr-Lindstrom H.; (In) Bach M.K. (eds.); Immediate hypersensitivity: modern concepts and developments, Marcel Dekker, New York (1978).
                                                                                                                   PRELIMINARY SEQUENCE (MYELOMA PRECINICH H.H., Johansson S.G.O.,
                                                                                                                                                                                                                 Flanagan J.G., Rabbitts T.H.;
"The sequence of a human immunoglobulin epsilon heavy chain constant region gene, and evidence for three non-allelic genes.";
EMBO J. 1:655-660(1982).
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. MEDLINE=84236029;
                                                                                                                                                                                                                                                                                                              Max B.E., Battey J., Ney R., "Duplication and deletion in Cell 29:691-699(1982).
                                                                                                                                                                                                                                                                                                                                                                      MEDIJINE=8316897; PubMed=6300763; Seno M., Kurokawa T., Ono Y., Onda H., Sasada R., Igar Kikuchi M., Sugino Y., Nishida Y., Honjo T.; "Molecular cloning and nucleotide sequencing of human epsilon chain cDNA."; Nucleic Acids Res. 11:719-726(1983).
                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.,
MEDLINE=83001945; Pt
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Eukaryota; Metazoa; C
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., von Bahr-Lindstrom H.;
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PDB; 1F95; X-ray; A=207-428.
PDB; 1G84; MNR; A=106-210.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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"A model of the Fc of immunoglobulin
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IGHE.
                                                                                                                                                                         PubMed=3796618; DOI=10.1016/0161-5890(86)90005-2;
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MEDLINB-84236092; PubMed-6329728;
Ishida N., Ueda S., Hayashida H., Miyata T.,
Ishida N., Ueda S., Hayashida H., Miyata T.,
"The nucleotide sequence of the mouse immunog
comparison with the human epsilon gene sequence EMBO J. 1:1117-1123(1982).
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                                                                                                                              SEQUENCE OF 34-421 FROM N.A.
MEDLINE=83117774; PubMed=681855;
Liu F.-T., Albrandt K., Succliffe
"Cloning and nucleotide sequence of
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3; Mismatches 6
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PIR; A02145; EHMSS.
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PROSITE; PS50835; IG LIKE;
PROSITE; PS00290; IG_MHC;
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rPro; IPR007110; 1
rPro; IPR003597; 1
rPro; IPR003006; 
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"A new antigen receptor gene family extensive somatic diversification Nature 374:168-173(1995).
EMBL; U18701; AAB48195.1; -.
PIR; S60266; S60226.
HSSP; P01842; 1AQK.
                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Cran
Blasmobranchii; Galeomorphii; Gale
Ginglymostomatidae; Ginglymostoma.
                                                                                                                                                                                                                                                                                                                                          01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NAR-2004 (TrEMBLrel. 26, Last annotation update)
Novel antigen receptor precursor
Ginglymostoma cirratum (Nurse shark)
                                                                                                                               Greenberg A.S., Flajník M.F.;
                                                                                                                                                                                                                                                       Ginglymostomatidae;
NCBI_TaxID=7801;
                                                                                                                                                                     TISSUE=Spleen;
MEDLINE=95183140;
                                                                                                                                                                                                                 EQUENCE FROM N.A.
                                                                                                                                                 0; PubMed=7877689;
Avila D., Hughes!
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Galeoidea;
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Noore T., Max S.I., Wang J., Hsieh F.K.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonamer A.A., Rubin G.M., Hong L.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA RAMAS S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schmerch A., Schein J.E.,
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Klein S., Strausberg R.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; BC072094; AAH72094.1; -.
                                                                                                                                SEQUENCE FROM N.A. TISSUE-Heart;
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Jones S.J., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                    Klein S.L., Strausberg Richardson P.;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Receptor; Signal.
SIGNAL 1
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                                                                                                                                                                                                                                                                                                                                                                        PubMed=12454917; DOI=10.1002/dvdy.10174; sberg R.L., Wagner L., Pontius J., Clifton S.W.,
                                                                                                                                                                                                                                                                                                                    tools
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75224 MW;
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Pred. No. 7
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novel antigen receptor.
, 2FF9D2071CDA6DFD CRC64;
                                                                                                                                                                                                                                                                                                Xenopus research: The NIH Xenopus
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RESULT 7
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ID MYH6
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Best Local S
Matches 10
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GO; GO:0016459; C:myosin; IEA.
GO; GO:0005524; F:MTO binding; IEA.
GO; GO:000374; F:motor activity; IE
InterPro; IPR001609; Myosin head.
Pfam; PF00063; Myosin head; 1.
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Best Local (
P02563; Q63351;
21-JUL-1986 (Rel. 01, Created)
01-CCT-1989 (Rel. 12, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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GO; GO:0016459; C:myosin; IEA.
GO; GO:000524; F:ATP binding; IEA.
GO; GO:0005774; F:motor activity; II
InterPro; IPR000048; IQ region.
InterPro; IPR001609; Myosin_Nead.
InterPro; IPR001609; Myosin_N.
Pfam; PF00612; IQ; 2.
Pfam; PF00062; Myosin_head; 1.
Pfam; PF00073; Myosin_Nead; 1.
PFINTS; PR00193; MYOSINHEAVY.
ProDom; PF000355; Myosin_head; 1.
SMART; SM00214; MYSc; II.
SMART; SM00215; IQ; 2.
SMART; SM00215; IQ; 2.
SMART; SM00215; IQ; 1.
NON TER 1080 Ab. 12024 MY.
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
STRAIN-New Zealand White; TISSUE-Heart;
STRAIN-New Zealand White; TISSUE-Heart;
Sayeed R.A., Grace A.A., Vandenberg J.I.;
Sayeed R.A., Grace A.A., Vandenberg J.I.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              09MYS5, PRELIMINARY; PRT; 99 AA.
09MYS5, CTEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence up
01-OCT-2003 (TrEMBLrel. 25, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                           Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Lagomorpha; Leporidae; Oryctolagus.
NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=myh6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Myosin heavy chain
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                                                                                                                                              RVTHPHLPKDIVRSIAKAPG 21
                                                                                                                           RTTHPHFVRCIIPNERKAPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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1080 AA;
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                                                                                                                                                                      Score 54; DB;
Pred. No. 5.5;
3; Mismatches
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                                                                                                                           98
                                                                                                                                                                                                                   3961CA1047DB2E0C CRC64;
                                                                  PRT;
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                                                                  1938
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46;
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(MyHC-alpha).

B

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ELEUT. Heart J. 5:181-191(1984).

C. --- SUBUNIT: Muscle contraction.

C. --- SUBUNIT: Muscle myosin is a hexameric protein that consists of 2 heavy chain subunits (MHC), 2 alkali light chain subunits (MLC).

C. --- SUBUNIT: Muscle myosin is a hexameric protein that consists of 2 heavy chain subunits (MHC), 2 alkali light chain subunits (MLC).

C. --- SUBURIT: Muscle myosin subunits (MLC-2).

and 2 regulatory light chain subunits (MLC-2).

--- SUBURITARIORATION: Thick filaments of the myofibrils.

C. --- DOMAIN: The rodlike tail sequence is highly repetitive, showing cycles of a 28-residue repeat pattern composed of 4 heptapeptides, cycles of a 28-residue repeat pattern composed of 4 heptapeptides, cycles of a 28-residue repeat pattern composed of 4 heptapeptides, cycles of a 28-residue repeat pattern combosed of 1 ight meromyosin (LMM) and 1 heavy meromyosin (LMM). It can later be split further into 2 globular subfragments (S1) and 1 rod-shaped subfragments (S2).

C. --- MISCELLANBOUS: The cardiac alpha isoform is a 'fast' ATPase myosin, while the beta isoform is a 'slow' ATPase.

--- SIMILARITY: Contains 1 Myosin-like globular head domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in the
Nature
[5]
EMBL; X15938; CAA34064.1; -.
EMBL; X01464, AAA41648.1; -.
EMBL; J00751; AAA41653.1; -.
EMBL; M32697; AAA41658.1; -.
PIR; S06005; S06005.
                                                                                                                                                                                entities requires a license agreement (See http:\overline{/}/www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mahdavi V., Lompre A.M., Chambers A.P., Nadal-Ginard B., 
"Carddac myosin heavy chain isozymic transitions during development 
and under pathological conditions are regulated at the level of mRNA 
availability.",
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 1-167 FROM N.A.
MEDLINE-84194059; PubMed-6585819;
Mahdavi V., Chambers A.P., Nadal-Ginard B.;
"Cardiac alpha- and beta-myosin heavy chain tandem.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McNally E.M., Kräft R., Bravo-Zehnder M., Taylor D., Leinwand L.A.; "Pull-length rat alpha and beta cardiac myosin heavy chain sequences. Comparisons suggest a molecular basis for functional differences."; J. Mol. Biol. 210:665-671(1989).
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Kraft R., Bravo-Zehnder M., Taylor D., Leinwand L.A.;
"Complete nucleotide sequence of full length cDNA for rat alpha cardiac myosin heavy chain.";
"Cardiac myosin heavy chain.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=Myh6;
Rattus norvegicus (Rat).
Rukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-Wistar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mahdavi V., Periasamy M., Nadal-Ginard B.; "Molecular characterization of two myosin
                                                                                                                                                                                                                                                     modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=85179510; PubMed=6241892;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 1872-1938 FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adult heart.";
297:659-664(1982)
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Am. J. Med.
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InterPro; IPR002928;
Pfam; PF00612.
SEQUENCE FROM N.A.

MEDLINE-94140346; PubMed-8307559;

Epp T.A., Dixon I.M., Wang H.Y., Sole
"Structural organization of the human
gene (MYH6).";
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Pfam; PF02736; Myosin_N; 1.
Pfam; PF01576; Myosin_tail 1; 1.
PFINTS; PR00193; MYOSINHEAVY.
PTODOm; PD000355; MYOSINHEAVY.
PROSITE; PS50096; IQ; 1.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                        Matsuoka R., Beisel K.W., Fu
"Complete sequence of human
                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                  MEDLINE=92133665; PubMed=1776652;
Matsuoka R., Beisel K.W., Furutan
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                        comparison to other myos differences.";
. Genet. 41:537-547(1991)
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R -> AP (in Ref. 3).

V -> A (in Ref. 3).

VS -> AP (in Ref. 3).

B -> Q (in Ref. 3).

Missing (in Ref. 3).

P -> PF (in Ref. 4).

R -> S (in Ref. 4).

R -> T (in Ref. 4).

T -> N (in Ref. 4).
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                                       Sole M.J., Liew
human cardiac alp
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alpha-myosin heavy chain
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alpha-myosin
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family; Muscle Protein; Myosin; Thick filament

Page

Coiled coil (Potential).

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DR EMBL, DOS-13. ILCENSE agreement (See http://w
DR EMBL, DOS-13. BAAOO791.1,
DR EMBL, M25140; AAA79675.1;
DR EMBL, M25140; AAA60386.1;
DR EMBL, M25142; AAA60386.1;
DR EMBL, M25142; AAA60387.1;
DR EMBL, M25142; AAA60387.1;
DR EMBL, M2164; AAA60387.1;
DR HASP, P24733; IKQM.
DR GO, GO:000524; F. Calmoduling; NAS.
DR GO; GO:000524; F. Calmoduling; NAS.
DR GO; GO:000526; F. Calmoduling; NAS.
DR InterPro; IPR00049; IQ. zegion.
DR InterPro; IPR00049; Myosin. head.
DR Fam; PR00150; Myosin. head.
DR Fam; PR00150; Myosin. head.
DR Fam; PR00150; Myosin. head.; I.
PROMORDS; Myosin. head.; I.
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Proc. Natl. Acad. Sci. U.S.A. 86:7416-7417(1989). Liew C.-C.,
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**MEDLINE:=89264452; PubMed=2726733; PROM N.A.

**Characterization of human cardiac myosin heavy chain genes.", [4]

**Indian of human cardiac myosin heavy chain genes.", [4]
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                                                                                                                                                         SEQUENCE OF 1630-1939 FROM N.A.

MEDLINE-86205859; PulMed=3458174;

"Construction and characterization of the alpha form of a cardiac syrian heavy chain cDNA clone and its developmental expression in the construction and characterization of the alpha form of a cardiac syrian heavy chain cDNA clone and its developmental expression in the contraction 3175-3179(1986)
                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN.B1B; TISSUB=Liver;
MEDLINE.9511503; PubMed=7815459; DOI=10.1006/jmcc.1994.1134;
MEDLINE.9511503; PubMed=7815459; DOI=10.1006/jmcc.1994.1134;
MAIR R., Sole M.J., Cukerman E., Liew C.-C.,
Diany Chain 9ene from Syrian hamster.";
Mol. Cell. Cardiol. 26:1155-1165(1994).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mesocricetus auratus (Golden hamster).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia, Butheria; Rodentia, Sciurognathi; Muridae, Cricetinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P1533; PARTINIARU;

O1-UAN-1990 [Rel. 13, Created]
05-UUL-2004 [Rel. 37, Last sequence update]
Myosin heavy chain, cardiac muscle alpha isoform (MyHC-alpha).
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Actin-binding.

Actin-binding.

NG, NG, NG-crimeshyllysine (Potential)

Q = P (in Ref. 1).

A -> P (in Ref. 1).

Y -> A (in Ref. 1).

A -> Y (in Ref. 1).

Y -> A (in Ref. 1).

A -> Y (in Ref. 1).

A -> Y (in Ref. 1).

A -> Y (in Ref. 1).

Y -> NV (in Ref. 1).

Y -> NV (in Ref. 1).

Y -> NV (in Ref. 1).

Y -> Y (in Ref. 2).

Y -> Y (in Ref. 3).

Y -> Y (in Ref. 3).

Y -> Y (in Ref. 3).

Y -> Y (in Ref. 5).

Y -> Y (in Ref. 7).

Y -> Y (in Ref. 7).

Y -> Y (in Ref. 7).

Y -> Y (in Ref. 1).

Y -> Y (in Ref. 1).

Y -> Y (in Ref. 1).

Y -> Y (in Re
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50.0%; Pred. No. 1.20+02;
tive 3; Mismatches 7;
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Indels

0, Gaps

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RESULT 10
Q7RZJ3
ID Q7RZJ3
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                                                                                                                       Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                              Pfam; PP00612; IQ; 2.
Pfam; PP0063; Myosin_head; 1.
Pfam; PP02736; Myosin_tail_1; 1.
Pfam; PP01576; Myosin_tail_1; 1.
Pfam; PP010315; MYOSINHEAVY.
PRODOM; PD000355; MYOSINHEAVY.
PRODOM; PS00096; IQ; 1.
PROSITE; PS50096; IQ; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000048; I
InterPro; IPR001609; N
InterPro; IPR004009; N
InterPro; IPR002928; N
                                                                                                                                                                                                     MOD RES
CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; L15351; AAB59701.1; -.
EMBL; M12995; AAA37081.1; -.
PIR; 148175; 148175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Actin-binding; ATP-binding; Calmodulin-binding; Coiled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSSP; P24733; 1KQM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBUNIT: Muscle myosin is a hexameric protein that consists of 2 heavy chain subunits (MHC), 2 alkali light chain subunits (MLC) and 2 regulatory light chain subunits (MLC-2).

SUBCELLULAR LOCATION: Thick filaments of the myofibrils.

DOMAIN: The rodlike tail sequence is highly repetitive, showing cycles of a 28-residue repeat pattern composed of 4 heptapeptides, characteristic for alpha-helical coiled coils.

MISCELLANEOUS: Each myosin heavy chain can be split into 1 light meromyosin (LMM) and 1 heavy meromyosin (HMM). It can later be split further into 2 globular subfragments (S1) and 1 rod-shaped
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MISCELLANEOUS: The cardiac alpha isoform is a 'fast' ATPase myosin, while the beta isoform is a 'slow' ATPase. SIMILARITY: Contains 1 IQ domain.
SIMILARITY: Contains 1 myosin-like globular head domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      subfragment (S2)
                                                                    665
                                                                                             N
                                                                                                                                     Similarity
                                                                                RVTHPHLPKDIVRSIAKAPG
                                                                    RTTHPHFVRCIIPNERKAPG
                                                                                                                                                                             1633
1651
1686
1693
1844
1879
1879
1907
1907
1928
1933
  PRELIMINARY;
                                                                                                                       Conservative
                                                                                                                                                                             Ŗ,
                                                                                                                                                             1935
AA; 223626 MW;
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1844
1879
1885
1907
                                                                                                                                                                                                                                                                                       1687
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Myosin_head.
Myosin_N.
Myosin_tail.
                                                                                                                                  39.7%;
                                                                                                                                                                       Q -> I (in Ref. 2).
H -> Q (in Ref. 2).
SL -> DV (in Ref. 2).
V -> G (in Ref. 2).
K -> R (in Ref. 2).
A -> T (in Ref. 2).
B -> Q (in Ref. 2).
B -> V (in Ref. 2).
D -> N (in Ref. 2).
D -> N (in Ref. 2).
D -> N (in Ref. 2).
C -> N (in Ref. 2).
                                                                                                                       ω
T
                                                                                                                                    Score 54;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                       Poly-Ala.
ATP (Potential).
Actin-binding.
                                                                                                                                                                                                                                                                                                                                                                                               Coiled coil
                                                                                                                                                                                                                                                                                                                                                                                                                          Myosin
                                                                                                                                                                                                                                                                                                                              N6,N6,N6-trimethyllysine (Potential)
                                                                                                                                                                                                                                                                                                                                          Actin-binding
  PRT;
                                                                    684
                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                           Muscle protein;
yosin head-like.
                                                                                                                          Mismatches
 538
                                                                                                             DB 1,
1.2e+02;
7;
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                                                                                                                                                                                                                                                                                                                                                                                                (Potential)
                                                                                                                                                  Length 1939;
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                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                         coil;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           filament.
                                                                                                                       Gaps
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RA Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B., Ra Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M., Qui D., Ianakiev P., Pedersen D., Nelson M., Washburne M., Ra Selitrennikoff C.P., Kinsey J.A., Braun E.L., Zelter A., Schulte U., RA Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D., Ramussen C., Marcotte E., Greenberg D., Ramal M., Kamyyselis M., Maucell E., Bielke C., Rudd S., Frishman D., Kantyseolis M., Maucell E., Bielke C., Rudd S., Frishman D., Kantyseolis M., Maucell E., Bielke C., Rudd S., Frishman D., Kochen S., Ra Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S., Ra Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S., DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R., DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R., Paulsen I., Sachs M.S., Lander S., Dunlap J., Radford A., Aramayo R., Paulsen I., Sachs M.S., Lander S., Nusbaum C., Birren B.; The Genome Sequence of the Filamentous Fungus Neurospora crassa."; RL Nature 0:0-0(2003):

CC EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is preliminary data.

SEQUENCE 538 AA; 61410 No.
                                                                                             RESULT
QREAD
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Best Local S
Matches 11
  Query Match
Best Local S
Matches 11
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Q870P3;
01-JUN-2003
01-JUN-2003
01-MAR-2004
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01-MAR-2004
01-MAR-2004
01-MAR-2004
                                                                                                                                              German Neurospora genome project; submitted (MAR-2003) to the EMBL/GenBank/DDBJ EMBL; BX295540; CAD79685.1; -.
                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Fungi; Ascomycota; Posordariomycetidae; Sordariales; NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Pungi; Ascomycota; Pe
Sordariomycetidae; Sordariales;
NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                     Schulte U., Aign V., Hoheisel J., Brandt P., Nyakatura G., Mewes H.W., Mannhaupt G.; Submitted (MAR-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                [1]
SEQUENCE FROM N.A.
Tr. Aign V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical protein Name=49D12.100;
                                                                                                   Hypothetical SEQUENCE 9(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neurospora crassa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neurospora crassa.
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                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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  ch 39.0%;
l Similarity 50.0%;
ll; Conservative
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, Last annotation update)
.100.
; Score 53; DB
; Pred. No. 76;
3; Mismatches
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Last sequence update)
Last annotation update)
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Pezizomycotina;
; Sordariaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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                                                                                                71EDE49AF8E03146
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                                                  DB 2;
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                                                                                                                                                                          databases
                                                                                                                                                                                                                                                                        databases
                                                                                                                                                                                                                                                                                                                                                                                                                   Sordariomycetes;
Neurospora.
                                                Length 906;
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                                                                                                   CRC64;
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RESULT Q60096
ID Q6
AC Q6
DT 05
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Best Local
EMBL; BX640620; CAB45774.1; -. HSSP; P01861; 1ADO.
InterPro; IPR003599; IG.
InterPro; IPR007110; IG-like.
InterPro; IPR003597; IG-c1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q6N096;
05-JUL-2004
05-JUL-2004
                                                                                                                                                                                    Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
Hypothetical protein DKFZp686I15196.
Name=DKFZp68GI15196;
                                                                                                                                                                                                                                                                  The German Human cDNA Consortium;
Wambutt R., Heubner D., Mewes H.W., Weil B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-ATCC 1908) / CB15;

STRAIN-ATCC 1908) / CB15;

MEDLINE-21173698; PubMed=11259647; DOI=10.1073/pnas.061029298;

Nierman W.C., Feldblyum T.V., Laub M.T., Paulsen I.T., Nelson K.E.,

Eisen J.A., Heidelberg J.F., Alley M.R.K., Ohta N., Maddock J.R.,

Potocka I., Nelson W.C., Newton A., Stephens C., Phadke N.D., Ely B.,

Potocka I., Nelson W.C., Newton A.S., Gwinn M.L., Haft D.H.,

DeBoy R.T., Dodson R.J., Durkin A.S., Gwinn M.L., Haft D.H.,

Kolonay J.F., Smit J., Craven M.B., Khouri H.M., Shetty J.,

Berry K.J., Utterback T.R., Tran K., Wolf A.M., Vamathevan J.J.,

Ermolaeva M.D., White O., Salzberg S.L., Venter J.C., Shapiro I.,
                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
TISSUE-Human esophagus tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q6N096
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Proc. Natl. Acad. Sci. U.S.A. 98:4136-4141(2001).
EMBL; AE005730; AAX22550.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIGR; CC0564; -.
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein CC0564.
OrderedLocusNames=CC0564;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9AAN3;
01-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Caulobacter crescentus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 THPHLP--KDIVRS-----IAKAPGKRAP 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SRPHAPRNRDIIRSTLGRDKPLCARANGRRAP 255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proteome; Hypothetical protein.
273 AA; 29461 MW; ClC6D2FB7E432082 CRC64;
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ilarity 40.6%;
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Primates;
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Pred. No. 31;
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Q6P8P3; 05-JUL-2004 05-JUL-2004

(TrEMBLrel. PRELIMINARY;

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Q6P8P3

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STRAIN-A3(2) / M.45;

STRAIN-A3(2) / M.45;

MEDLINE-21996410; PubMed=12000953; DOI=10.1038/417141a;

MEDLINE-21996410; PubMed=12000953; DOI=10.1038/417141a;

MEDLINE-21996410; PubMed=12000953; DOI=10.1038/417141a;

Bentley S.D. Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L.,

Thomson N.R., James K.D., Harris D.B., Quail M.A., Kieser H.,

Harper D., Bateman A., Brown S., Chandra G., Chen C.W., Collins M.,

Harper D., Bateman A., Brown S., Chandra G., Chen C.W., Collins M.,

Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,

Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,

Huang C.-H., Kieser T., Larke L., Murphy L.D., Oliver K., O'Neil S.,

Huang C.-H., Kieser T., Larke L., Murphy L.D., Oliver K., O'Neil S.,

Huang C.-H., Rieser R., Galandream M.A., Rutherford K.M., Rutter S.,

Taylor K.,
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InterPro; IPR003596; Ig_v.
Pfam; PP07654; C1-Bet; 3.
SMART; SM00409; IG; 2.
SMART; SM00407; IGC1; 3.
SMART; SM00406; IGV; 1.
SMORT; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_MHC; UNKNOWN_2.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
SEQUENCE 466 AA; 50926 MW; 01B91B7
                                                                                                                                                                                                                                      InterPro; IFR008938; ARM.
InterPro; IFR008938; PBS lyase_HEAT.
Pfam; PF03130; HEAT_BS; 1.
SMART; SM00567; EZ HEAT; 4.
Complete proteome; Hypothetical prot
SEQUENCE 197 AA; 21061 MM; 6ABC3
                                                                                                                                                                                                                                                                                                                                                "Complete genome sequence of tocolicolor A3(2).";
Nature 417:141-147(2002).
EMBL, AL939127; CAA22724.1; -...
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Seeger K., Saunders D., Sharp S., Squares R., Squares S.,
Warren T., Wietzorrek A., Woodward J.R., Barrell B.G., Par
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9ZBJ8, PRELIMINARY,
Q9ZBJ8, 01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                     PIR; T35963; T35963
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Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomyces.
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                                                                                                                                                                                Similarity 9; Conserv
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Pred. No.
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Last annotation updat
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Pred. No. 63;
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6ABC3643C5570AC1 CRC64;
                                                                                                                    197
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RP SEQUENCE FROM N.A.

RC TISSUB-Lung and heart;

RC TISSUB-Lung and heart;

RC TISSUB-Lung and heart;

RM Strausborg R.L., Peingold B.A., Grouse L.H., Derge J.G.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Mang J., Hsieh F.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.K.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.B.,

RA Stapleton M.J., Uddin T.B., Toshiyuki S., Carninoi P., Prange C.,

RA Stapleton M.J., Uddin T.B., Toshiyuki S., Carninoi P., Prange C.,

RA Brownstein M.J., Uddin T.B., Toshiyuki S., Carninoi P., Prange C.,

RA Raha S., Loquellano N.A., Peters G.J., Abzamson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Say L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Say L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Solbbs R.A.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Solbbs R.A.,

RA Richards S., Worley K.C., Shevchenko Y., Bouffard G.G.,

RA Rabey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Mitting M., Madan A., Young A.C., Shevchenko Y., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Jones S.J., Warra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

RT and mouse cDNA sequences.";

RT Total Radia A. S., Scherch A., Scherch A., Schein J.E.,

RA Krzywinski M.L., Acad. Sci. U.S.A. 99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

C TISSUE=Lung and heart;

Strausberg R.;

A Strausberg R.;

A Submitted (NOV-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; BC061145; AAH61145.1; -.

R EMBL; BC061145; AAH61145.1; -.

R GO; GO:0007522; P:visceral muscle development; IMP.

R GO; GO:0007522; P:visceral muscle development; IMP.

R InterPro; IPR0010048; IQ_region.

R InterPro; IPR00109; Myosin_Nead.

R InterPro; IPR004009; Myosin_N.

R Pfam; PF00061; IQ; 2.

R Pfam; PF00063; Myosin_Nead; 1.

Pfam; PF00063; Myosin_Nead; 1.
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Best Local Similarity
Matches 9; Conserva
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ProDom, PD000355; Myosin_head; 1.
SMART, SM00242; MYSC; 1.
PROSITE; PS50096; 1.
PROSITE; PS50096; 1.
NON TEE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-2004 (TrEMBLrel. 27, Last annotation update) Myh6 protein (Fragment). Name=Myh6;
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45.0%; Pred. No. 1.7e+02;
tive 4; Mismatches 7
                                 2005, 07:00:16
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protein search, using sw model
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Copyright (c) 1993 - 2005 Compugen Ltd.
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                             ribosomal protein bradykinin-potein bradykinin-potenti superoxide dismuta agrin - electric r bombolitin IV - Am flagellar motor swribosomal protein glutamate-ammonia kinetoplast DNA-as hypothetical prote somatotropin - Atl bradykinin-potenti outer layer protei retinoid-x-recepto
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acid ribonuclease
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granulocyte inhibi
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	Query Matc Best Local Matches	RESULT 2 878765 878765 C;Species: Bos prim C;Date: 20-Sep-1999 C;Accession: 878765 R;Graack, H.R. submitted to the pr A;Reference number: A;Accession: 878765 A;Molecule type: pr A;Residues: 1-11 <g a;cross-references:="" c;keywords:="" f;1-11="" mitchon="" product:="" rib<="" td=""><td>Best Local Matches</td><td>Eur. J. Blochem. 187 A;Title: Core histon A;Reference number: A;Accession: S07770 A;Molecule type: pro A;Residues: 1-25 <hi. a;cross-references:="" c;keywords:="" chromoso<="" td=""><td>RESULT 1 S07770 histone H2B.1, sperm C;Species: Echinus er C;Date: 30-Sep-1991: C;Accession: S07770 R;Hill, C.S.; Thomas</td><td>0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0</td></hi.></td></g>	Best Local Matches	Eur. J. Blochem. 187 A;Title: Core histon A;Reference number: A;Accession: S07770 A;Molecule type: pro A;Residues: 1-25 <hi. a;cross-references:="" c;keywords:="" chromoso<="" td=""><td>RESULT 1 S07770 histone H2B.1, sperm C;Species: Echinus er C;Date: 30-Sep-1991: C;Accession: S07770 R;Hill, C.S.; Thomas</td><td>0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0</td></hi.>	RESULT 1 S07770 histone H2B.1, sperm C;Species: Echinus er C;Date: 30-Sep-1991: C;Accession: S07770 R;Hill, C.S.; Thomas	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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HPHLPKDIVR : : : : HVDVPKDLTK :	h Similarity 4; Conservat	RESULT 2 878765 878765 878765 C;Species: Bos primigenius taurus C;Date: 20-Sep-1999 #sequence_rev C;Accession: 878765 R;Graack, H.R. submitted to the Protein Sequence A;Reference number: 878760 A;Accession: 878765 A;Molecule type: protein A;Residues: 1-11 <gra> A;Cross-references: UNIPROT:Q7M37 C;Keywords: mitochondrion F;1-11/Product: ribosomal protein</gra>	Similarity 5; Conservat KAPGKRAP 25 : : KSPTKRSP 11	Bur. J. Blochem. 187, 145-153, 19 A; Tattle: Core histone-DNA interace A; Reference number: 807769; MUID: A; Recession: 807770 A; Molecule type: protein A; Residues: 1-25 <hil- a;="" c;="" chromosomal="" cross-references:="" h2b="" histone="" keywords:="" march<="" oner:="" p1328="" protein;="" superfamily:="" td="" uniprot:=""><td>Echinus esculentus Sep-1991 #sequence Sep-1991 #sequence</td><td>117.3 117.3 116.9 116.9 116.9 116.9 116.9 116.9</td></hil->	Echinus esculentus Sep-1991 #sequence Sep-1991 #sequence	117.3 117.3 116.9 116.9 116.9 116.9 116.9 116.9
14 11		y misseque	62.5%; tive	-153 int 9; M	Bea urchin lentus quence_rev	222 233 240 250 250 250 250 250 250 250 250 250 25
	20.6%; 10.0%;	mitoch taurus nce_rev equence equence	65 40 14	, 19 erac UID: UID: 1328	rev	00000000000000000000000000000000000000
	Score 28; DB 2; L Pred. No. 7.9e+02; 3; Mismatches 3;	ondrial - bovine (fr (cattle) (sion 20-Sep-1999 #t Database, July 1999	9+03; 1	90 tions in sea urchin 90126812; PMID:2298 1 1 DNA binding; nucleo	ALIGNMENTS - sea urchin (Echinus esculentus) (sculentus #sequence_revision 30-Sep-1991 #text	942567 PS0273 A41263 A71263 B371990 B37109 PH1317 B27504 PA0012 PA0012 PA012 PA015 P
	Length; Inde	(fragment) #text_cha. 999	ğ '	sperm 202 some co	×	
	gth 11; Indels	ınge	Indels	chro	(fragment t_change	cytochrome-b5 hypothetical; kinase-relate cytochrome-c cytochrome-c hypothetical Ig heavy chai major allerge histone H2B - superoxide di carboxylester Ca2+/calmodul T cell recept ATPase R1 sub stromelysin (c probable acr- platelet aggr
	0,	ge 09-Jul-	0,	matin. nucleus	09-Jul-	ochrom ochrom ochrheti ochrom ochheti heavy or all trone H
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	0,	4 A	0;	N-terminal	•	ដើល់ធំធី `` ភិបិថា សំរ ៈ . ល្ម

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RESULT 3

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gene J protein - phage G4
C;Species: phage G4
C;Species: phage G4
C;Date: 30-Nov-1979 #sequence_revision 30-Nov-1979 #text_change 09-Jul-2004
C;Accession: A04259
R;Godson, G.N.; Barrell, B.G.; Staden, R.; Fiddes, J.C.
Nature 276, 236-247, 1978
A;Title: Nucleotide sequence of bacteriophage G4 DNA.
A;Reference number: A03200; MUID:79053264; PMID:714153
A;Accession: A04259
A,Molecula trace
                                                                                                                                A;Cross-references: UNIPROT:P03652; GB:J02454; GB:M10724; GB:M11404; GB:V00657; NID:g156
C;Comment: Gene J protein is one of the structural components of the bacteriophage coat
                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-25 < GOD>
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J. Blochem. 103, 267-273, 1988
A;Title: Purification of acid ribonucleases from bovine
A;Reference number: A41439; MUID:88227899; PMID:313131316
A;Accession: A41439
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C;Species: Bos primigenius taurus (cattle)
C;Date: 21-Apr.1992 #sequence_revision 21-Apr-1992 #text_change
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A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1351
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Ig heavy chain DJ region (clone C100-109B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1351
C;Accession: PH1351
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C;Keywords: heterotetramer; immunoglobulin
                                                                                                       Superfamily: phage phi-X174 gene J protein Keywords: DNA binding
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A; Residues: 1-16 < WAS>
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Best Local
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                               Score 27; DB
Pred. No. 2.5e
2; Mismatches
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Pred. No. 2e+03;
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2.5e+03;
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glutathione transferase (BC 2.5.1.18) mu (isoform pI 6.4) - mouse (fragment C;Species: Mus musculus (house mouse)
C;Date: 02-Dec-1993 #sequence_revision 01-Dec-1995 #text_change 09-Jul-2004
C;Accession: 832551; 832550, M.; Ahmad, H.; Awasthi, Y.C.
R;Singhal, S.S.; Baxena, M.; Ahmad, H.; Awasthi, Y.C.
Biochim. Biophys. Acta 1116, 137-146, 1992
A;Title: Glutathione S-transferases of mouse liver: sex-related differences
A;Reference number: 832548; MUID:92256466; PMID:1581342
A;Reference number: 332548; MUID:9;
A;Accession: 332551
A;Molecule type: protein
A;Residues: 1-13 <SINI>
A;Cross-references: UNIPROT:Q7M059
A;Experimental source: female
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Comment: This protein is specifically translated in the pollens. C;Comment: This protein functions by depolymerizing pectin in the y growing tube.
C;Keywords: glycosidase; hydrolase; nolveachers.
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A;Experimental source: pollen
C;Comment: This protein is specific
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A;Reference number: JQ0992; MUID:93005658; PMID:2152116
A;Accession: PQ0143
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A; Residues: 1-22 < BRO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Brown, S.M., Crouch, M.L.
Plant Cell 2, 263-274, 1990
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A;Molecule type: protein
A;Residues: 1-20 <HOB>
A;Cross-references: UNIPROT:Q7M4S4
C;Superfamily: immunoglobulin V region; immunoglobulin homology
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A;Title: hPysicochemical characterization of a polypeptide present A;Reference number: A36016; MUID:90349614; PMID:2385596
A;Accession: A36016
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A36016
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C;Species: Homo sapiens (man)
C;Date: 11-Jan-1991 #sequence_revision 11-Jan-1991 #text_change
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8; Conserv
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Pred. No. 2.
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                                                                                                                                      differences
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A; Molecule type: protein
A; Residues: 1-20 < CCT+A; Residues: 1-20 < CCT+A; Residues: 1-20 < CCT+A; Residues: UNIPROT: Q7M163
C; Superfamily: Escherichia coli ribosomal protein L30
C; Keywords: protein biosynthesis; ribosome
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JP0055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypothetical TEL/MN1 mutant fusion protein type I - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 20-Apr-2000
C;Accession: 138336
R;Buijs, A.; Sherr, S.; van Baal, S.; van Bezouw, S.; van der Plas, D.; Van Kessel, Oncogene 10, 1511-1519, 1995
A;Title: Translocation (12;22) (p13;q11) in myeloproliferative disorders results in A;Reference number: 138031; MUID:95249265; PMID:7731705
A;Accession: 138336
RESULT
H30608
                                                                                                                                                                                                                                                                                                                                            ribosomal protein L30 - Bacillus polymyxa (fragment)
C;Species: Bacillus polymyxa
C;Date: 10-Mar-1994 #sequence_revision 28-Oct-1994 #text_change 09-Jul-2004
C;Accession: JP0055
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A/Molecule type: protein
A/Residues: 1-13 <SIN2>
A/Experimental source: male
C;Keywords: transferase
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A;Description: Phylogenetic diversity
A;Reference number: JP0042
A;Accession: JP0055
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C; Keywords: fus:
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Map position: 22q11/12p13
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Best Local
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Pred. No. 1.8e+03;
                                                                                                                                          Score 26; DB 2;
Pred. No. 2.8e+03;
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Pred. No. 2.1e+03;
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J. Immunol. 142, 3158-3163, 1989
A;Title: Structural and idiotypic characterization of the L chains of human A;Reference number: A30601; MUID:89215279; PMID:2496160,
A;Accession: B30609
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R;Gon1, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson, J. Immunol. 142, 3158-3163, 1989
A;Title: Structural and idiotypic characterization of the L chains of human IgM A;Reference number: A30601; MUID:89215279; PMID:2496160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:Q9UL83; UNIPROT:Q9UL85
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig kappa chain V-III regions (Jon and Mit) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 09-Jul-2004
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D30609
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Reference number: A30601; MUID:89215279; PMID:2496160
A;Accession: H30608
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C;Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 09-Jul-2004
C;Accession: H30608
                                                                                       C; Superfamily: immunoglobulin V region; immunoglobulin C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                  C;Species: Homo sapiens (man)
C;Date: 29-Jun-1989 #sequence_revision 29-Jun-1989 #text_change 09-Jul-2004
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                                                                                                            A;Cross-references: UNIPROT:Q9UL83
C;Superfamily: immunoglobulin V region;
                                                                                                                                                       A; Molecule type: protein A; Residues: 1-24 < GON>
                                                                                                                                                                                                   A;Status: preliminary
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R;Goni, F.R.; Chen, P.P.; McGinnis, D.; Arjonilla, M.L.; Fernandez, J.; Carson,
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A; Residues: 1-22 < GON>
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4; Conservative
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Similarity 40.0%;
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Pred. No. 3.3e+03;
5; Mismatches 1
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Pred. No. 3e+03;
5; Mismatches
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bradykinin-potentiating peptide 8 - island jararaca
C;Species: Bothrops insularis (island jararaca)
C;Date: 14-Feb-1992 #sequence_revision 01-Dec-1992 #text_change 09-Jul-2004
C;Accession: H37196
R;Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
J. Protein Chem. 9, 221-227, 1990
A;Title: Primary structure and biological activity of bradykinin potentiating peptides in A;Reference number: A37196; MUID:90351557; PMID:2386615
A;Scatus: preliminary
A;Accession: H37196
A,Scatus: preliminary
A;Rose-references: UNIPROT:P30426
C;Keywords: pyroglutamic acid
C;Keywords: pyroglutamic acid
C;Keywords: pyroglutamic acid
C;Keywords: pyroglutamic acid
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H37196
Search completed: February 22, 2005, 07:20:44 Job time : 40 secs
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A; Description: Phylogenetic diversity in the genus Bacillus and comparative ribosomal
A; Reference number: JP0042
A; Raccession: JP0052
A; Molecule type: protein
A; Residues: 1-24 < COCH>
A; Cross-references: UNIPROT:07M162
C; Superfamily: Escherichia coli ribosomal protein L30
C; Keywords: protein biosynthesis; ribosome
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JP0052
CFDecies: Bacillus macquariensis (fragment)
CFSpecies: Bacillus macquariensis
CFDate: 10-Mar-1994 #sequence_revision 28-Oct-1994 #text_change 09-Jul-2004
CFAccession: JP0052
CFAccession: JP0052
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Best Local Similarity 50.0%;
Matches 5; Conservative
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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  seg length: 0 seg length: 25
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Match
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Gapop 10.0 ,
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  GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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canis famil
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hrassica ra
hacteriopha
plasmodium
escherichia
trypanosoma
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8 borrelia af
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Megener K.L., Wabnitz P.A., Carver J.A., Bowie J.H., Chia B.C.S.,
Wallace J.C., Tyler M.J.;
"Host defence peptides from the skin glands of the Australian blue
mountains tree-frog Litoria citropa. Solution structure of the
antibacterial peptide citropin 1.1.";
Eur. J. Biochem. 265:627-637(1999).
-I- SUBCELIULAR LOCATION: Secreted.
-I- TISSUE SPECIFICITY: Dorsal and submental skin glands.
Amphibian defense peptide; Direct protein sequencing.
SEQUENCE 18 AA; 1814 MW; 500BF778D515ABD7 CRC64;
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Submitted (SEP-2003) to the EMBL/GenBank/DDBJ
EMBL, AY39241; AAQ93549.1; -.
Hypothetical protein; Plasmid.
SEQUENCE 23 AA; 2374 MW; 6F371429BF49D3F8
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05-JUL-2004 (Rel. 44, Last annotation
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
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Republic of Congo.";
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MEDLINS=21195339; PubMed=11297743; DOI=10.1016/S0014-5793(01)02318-3;

HOLZMANN K., Ambrosch I., Elbling L., Micksche M., Berger W.;

"A small upstream open reading frame causes inhibition of human major vault protein expression from a ubiquitous mRNA splice variant.";

FEBS Lett. 494:99-104(2001).

EMBL; AJ291367; CAG35315.1; -.

Hypothetical protein.

Hypothetical protein.

SEQUENCE 18 AA; 2179 MW; 5D06F9A3F11CB828 CRC64;
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SEQUENCE.
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TISSUE-Skin secretion;
TISSUE-Skin secretion;
MEDLINE=99435977; PubMed=10504394;
Wegener K.L., Wabnitz P.A., Carver J
Wallace J.C., Tyler M.J.;
                                                                                                                                                                                                                                                                                                           P81838;
30-MAY-2000 (Rel. 39, C
30-MAY-2000 (Rel. 39, L
05-JUL-2004 (Rel. 44, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE:

MEDIINE-94198758; PubMeca:,,

Braun H.P., Kruft V., Schmitz U.K.;

Braun 193:99-106(1994).

Planta 193:99-106(1994).

GO; GO:0008121; F:ubiquinol-cytochrome-c reductase activity;

GO; GO:0008121; F:ubiquinol-cytochrome-c reductase activity;
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01-MAY-2000
01-MAY-2000
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01-MAY-2000 (TYEMBLYel. 13, Last sequence update)
01-CCT-2003 (TYEMBLYel. 25, Last annotation update)
Cytochrome-C reductase 14 kDa subunit (EC 1.10.2.2)
                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Anura; Ne
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                      Citropin 1.1.3.
Litoria citropa (Australian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embry
Spermatophyta; Magnoliophyta; eudicotyledona;
lamiida; Solanalea; Solanaceae; Solanum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Strausberg R.;
Submitted (APR-2001)
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                                                                                                                                                                              NCBI_TaxID=94770;
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Pred. No. 3.6e
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Pred. No. 4.1e+03;
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eobatrachia; Hyloidea; Hylidae;
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edons; core ev
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25-OCT-2004
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Q16271;
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Bur. J. Biochem. 265:627-637(1999).
-1- SUBCELLULAR LOCATION: Secreted.
-1- TISSUE SPECIFICITY: Dorsal and submental skin glands.
Amphibian defense peptide; Direct protein sequencing.
SEQUENCE 18 AA; 1814 MW; 500BF778D51F98D7 CRC64;
Ramachandran V., McArthur J.D., Behm C.E., Gutzeit C., Dowton M., Fagan P.K., Towers R.J., Currie B.J., Sriprakash K.S., Walker M.J.; "Distribution and molecular evolution of the fibronectin binding protein prtp2 gene family in Streptococcus pyogenes isolated from th Northern Territory of Australia.";
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MEDIJNS=95086168; PubMed=7993996;

Holtzman B.J., Kolakowski L.F.Jr., Geifman-Holtzman Rasoulpour M., Guillot A.F., Ausiello D.A.;

"Mutations in the vasopressin V2 receptor gene in two processins of the vasopressin v2 receptor gene in two processins v2 receptor gene in two p
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Eukaryota; Metazoa; C
Mammalia; Eutheria; P
                                                                                                                                                                                                                                                                                     Streptococcus.
NCBI_TaxID=1314;
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01-NOV-1996 (TYEMBLrel. 01, Last sequence update)
01-JUN-2003 (TYEMBLrel. 24, Last annotation update)
Arginine vasopressin V2 receptor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                   Streptococcus pyogenes.
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J. Am. Soc. Nephrol. 5:169-176(1994).
                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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1905 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21.3%;
                                                                                                                                                                                                                                                                                                                                                                                               Lactobacillales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8, Last sequence update)
8, Last annotation update)
regulator (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                181640EFD90F2788 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 4.8e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 4.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29;
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                                                                                                                                                                                                                                                                                                                                                                                               Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.6e+03;
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DRIAN OCC OCC DITION
                                                                                                                                                                                    RESULT 12
Q7M374
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H2B1_ECHES
                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                       01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Ribosomal protein MRP-S24, mitochondrial (Fragment).
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata: E
Graack H.R.;
Submitted (JUL-1999)
PIR; S78765; S78765.
                                                                                                                                                  Q7M374;
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H2B1 ECHES
P13281;
01-JAN-1990
01-JAN-1990
05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIINE=90126812; PubMed=2298202;
Hill C.S., Thomas J.O.,
"Core histone-DNA interactions in sea urchin sperm
terminal tail of H2B interacts with linker DNA.";
Eur. J. Biochem. 187:145-153(1990).
-i- SUBUNIT: The nucleosome is an octamer containin
each of H2A, H2B, H3 and H4. The octamer wraps
                                                                                                                                                                                                                                                                                                                                    PIR; S07770; S07770.

InterPro; IPR000558; Histone H2B.
PROSITE; PS00357; HISTONE H2B; PARTIAL.
Chromosomal protein; Direct protein sequencing; DNA-binding; Multigene family; Nuclear protein; Nucleosome core.

NON_TER 25
                                   SEQUENCE.
                                                       NCBI_TaxID=9913;
                                                                    Bovinae; Bos.
                                                                             Mammalia;
                                                                                                                                                                          Q7M374
                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Echinus esculentus (Sea urchin).
Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
Echinoidea; Euechinoidea; Echinacea; Echinoida; Echinidae; Echinus.
NCBI_TaxID=7648;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JAN-1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Histone H2B.1, sperm (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (JUL-2004) to the EMBL; AJ784873; CA404963.1; NON TER 21 SEQÜENCE 21 AA; 2332 MW;
                                                                                                                                                                                                                                                                                                                                                                                                      - SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             bp of DNA.
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                                                                                                                                                                                                                                                                            Similarity 5; Conserv
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|| | : ||::
5 PHFPINNVRNL 15
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                                                                                                                                                                                                                                                      KAPGKRAP 25
                                                                                                                                                                                                                               KSPTKRSP 11
                                                                               Eutheria;
                                                                                                                                                                                                                                                                                                                         25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 AA; 2332 MW;
                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                         2693 MW;
                                                                             Chordata; Craniat Cetartiodactyla;
          ç
                                                                                                                                                                                                                                                                                     21.3%;
62.5%;
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          the
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o the histone
          PIR
                                                                                                                                                                                                                                                                          Score 29; DB
Pred. No. 6.4e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                     9842DD3D73A3A9EC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
Pred.
                                                                           Craniata; Vertebrata;
actyla; Ruminantia; Pec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2086BDCA45FE51A1 CRC64;
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          data bank
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                                                                                                                                                                                                                                                                                                                                                                                                                                   octamer containing two
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                                                                                                                                                                                                                                                                                       6.4e+03;
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                                                                                                                                                                                                                                                                                                 DB 1;
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л.4е+03;
3;
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                                                                                                                                                                                                                                                                                              Length 25;
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                                                                            Pecora;
                                                                                                                                                                                                                                                                           Indels
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                                                                         Buteleostomi;
cora; Bovidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  molecules
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RESULT 14
Q9L7N8
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Q9MX47

ID Q9MX4A

AC Q9MX4

AC Q1

CO MCC ACCL

CO 
      SO TO REPORT OF THE PROPERTY O
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Best Local S
Matches 5
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Best Local (
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O9L7N8;
01-OCT-2000 (TrEMBLrel. 1
01-OCT-2000 (TrEMBLrel. 1
01-OCT-2000 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9MX47;
01-OCT-2000
01-OCT-2000
                                                                                  DOI=10.1128/JB.182.7.2037-2042.2000;
Gorbacheva V.Y., Godfrey H.P., Cabello F.C.;
"Analysis of the bmp gene family in Borrelia
J. Bacteriol. 182:2037-2042(2000).
                                   NON
                                                            EMBL;
                                                                                                                                                                                                                                                                                                                               Borrelia afzelii.
Bacteria; Spirochaetes;
NCBI_TaxID=29518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON_TER
NON_TER
SEQUENCE
      SEQUENCE
                                                                                                                                                                                                        STRAIN=Ip3;
MEDLINE=20179823; PubMed=10715014;
                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Name=bmpD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        BmpD (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Naruse K., Oku H., Kojima A.,
Hori H., Shima A., Nonaka M.,
Submitted (OCT-1999) to the EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryzias latipės (Medaka fish) (Japanese ricefish).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Atherinomorpha;
Beloniformes; Adrianichthyidae; Oryzinae; Oryzias.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MHC class II antigen (Fragment).
Name=Orla-DCB;
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2; Mismatches
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P82619;
16-OCT-2001
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MEDLINE-20189894, PubMed=10723010;

Predel R., Eckert M.;

Predel R., Eckert M.;

"Tagma-specific distribution of FXPRLamides in the nervous system of the American cockroach.";

J. Comp. Neurol. 419:352-363(2000).

-I- FUNCTION: Mediates visceral muscle contractile activity (myotropic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
05-UU-2004 (Rel. 44, Last annotation update)
Pyrokinin-4 (Pea-PK-4) (YXPRL-amide)
Periplaneta americana (American cockroach)
Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Orthopteroidea, Dictyoptera, Blattaria, Blattoidea,
Blattidae, Periplaneta.
                                                                                                                                                                                               -I- SIMILARITY: Belongs to the pyrokinin family.
Amidation; Direct protein sequencing; Neuropeptide; Pyrokinin.
MOD_RES 12 12 Leucine amide.
SEQUENCE 12 AA; 1449 MW; PA7A3049FP42CAA1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE, FUNCTION, AND MASS SPECTROMETRY.
TISSUB-Retrocerebral complex;
TISSUB-Retrocerebral complex;
MEDLINB-99212469; PubMed=10196736; DOI=10.1016/S0965-1748(98)00117-9;
Predel R., Kellner R., Nachman R.J., Holman G.M., Rapus J., Gaede G.;
"Differential distribution of pyrokinin-isoforms in cerebral and abdominal neurohemal organs of the American cockroach.";
Insect Biochem. Mol. Biol. 29:139-144(1999).
                                                                                                                                                                                                                                                                                  activity).
-i- TISSUE SPECIFICITY: Corpora cardiaca.
-i- MASS SPECIROMETRY: WW-1147.9; METHOD-MALDI; RANGE=1-12;
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RESULT 1 AAW24102 WPI; 1997-389423/36. N-PSDB; AAT85651. 30-JUN-1997. Canis familiaris. Canine immunoglobin B peptide AAW24102 standard; peptide; JP09169795-A. 21-NOV-1997 22-DEC-1995; 22-DEC-1995; Immunoglobulin E; IgE; anti-canine IgE antibody; allergy; canine; dog AAW24102; (HITB) HITACHI CHEM 95JP-00334381 (first entry) 95JP-00334381 CO LTD 20 ŗ ₿

Adq76006 Aao18028 Abj00529 Aao18018 Aao18041 Adg17011

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75.0%;

Score 102; Pred. No.

1.7e-08; DB 2;

Length 20, Indels

Aay68602 Aay91212 Aay79998 Aay80077 Aay80077

Optimised

Sequence 20 AA;

canine allergies

Aay50896 Adq76005 Aay50895 Adq76004 Aay80000 Aay50898 Adq76007

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AAW24098-106 are peptide fragments containing at least 5 continuous aminacids of the partial canine immunoglobulin E (IgE) protein shown in AAW24097. The peptides are used for the preparation of anti-canine IgE antibody. The anti-canine IgE antibody can be used for the diagnosis of

amino

Canine immunoglobulin ${\tt E}$ peptide fragment and related DNA - useful for the preparation of anti-canine immunoglobulin ${\tt E}$ antibody.

Claim 2, Page 9; 12pp; Japanese.

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RESULT 2
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                                                                                                                                                                                       Antibody 15A.2 binding peptide 10 from PhDc7c phage display library.
                                                                                                                                                                                                                                                                    24-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                 AAY50893 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel binding protein which specifically binds to native canine free or B-cell bound IgE, and which doesn't bind to IgE when the IgE is bound to mast cells. The peptide products of the invention have anti-allergic activity. The antibodies bind to defined epitopes on free or B-cell bound IgE molecules which have an important produce a pharmaceutical composition, preferably with a diluent, which can be used for prophylaxis or treatment of canine allergy. ANYS0876.

Y50900 represent peptide mimotopes used in the method of the invention
                                                                                                                                                                                                                                                                                                                          AAY50893,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding proteins used for treatment or prophylaxis of canine allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 7; 30pp; English.
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30-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody 15A.2 canine IgE binding epitope 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY50894 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-FEB-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 CRVTHPHLPKDIVRSI 16
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99US-00281760.
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                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 89; Pred. No.
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Best Local Similarity 100.
                   Claim 5; SEQ ID NO 2;
                                                New isolated polypeptides and encoding polynucleotides, useful for preventing and/or treating allergic diseases and allergy symptoms:
                                                                                       WPI; 2004-534123/51.
                                                                                                               Krah ER,
                                                                                                                                                          20-DEC-2002, 2002US-00325375.
                                                                                                                                                                                                                                                                                              Peptide capable of preventing IgE binding
                                                                                                                                                                                19-DEC-2003; 2003WO-US040711.
                                                                                                                                                                                                          15-JUL-2004.
                                                                                                                                                                                                                               WO2004058799-A2
                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                         allergy; allergic symptom; IgE binding
                                                                                                                                                                                                                                                                                                                         07-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel binding protein which specifically binds to native canine free or B-cell bound IgE, and which doesn't bind to IgE when the IgE is bound to mast cells. The peptide products of the invention have anti-allergic activity. The antibodies bind to defined epitopes on free or B-cell bound IgE molecules which have an important role in allergic reaction. The specific binding proteins are used to produce a pharmaceutical composition, preferably with a diluent, which can be used for prophylaxis or treatment of canine allergy. ANY50876.
                                                                                                                                      (IDEX-)
                                                                                                                                                                                                                                                                                                                                                  ADQ76003;
                                                                                                                                                                                                                                                                                                                                                                   ADQ76003 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Binding proteins used for treatment or prophylaxis of canine allergy.
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30-MAR-1999;
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                                                                                                                                    IDEXX LAB INC
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                                                                                                             Lawton R,
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99US-00281760.
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100.0%; Pred. No. 1.
tive 0; Mismatches
                  50pp; English.
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                                                                                                                                                                                                                                                                                              to high affinity receptor #2.
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The present invention

relates to isolated polypeptides capable

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inhibiting the binding of IgE to a high affinity receptor. These are useful for the prevention and/or treatment of allergic diseases and allergy symptoms in mammals. The present sequence is a polypeptide of the

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Matches 16
                        preventing triggering and activation of mast cells and basophils and downregulation of IgE synthesis. Conjugates, or fusion peptides, containing (I) are used for active immunisation against IgE-mediated allergies, e.g. food allergies, asthma, anaphylaxis, or flea-allergy dermatitis. Nucleic acids that encode these compounds are useful for recombinant production of corresponding peptides or in DNA vaccines. Conjugates of (I) that include a promiseuous T helper cell epitope (functional in genetically diverse subjects) in addition to a B cell target epitope, have increased immunogenicity and may include cyclic constraints (disulfide bridge) to stabilise conformational features and maximize cross-reactivity to the natural target. They induce safe (non-anaphylactogenic) antibodies. AAY79994 to AAY80084 represent amino acid sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunogenic; immunostimulatory; carrier protein; helper T cell epitope; antibody; allergy; allergic disease; immunisation; anti-allergic; anti-antibody; allergy; allergic disease; immunisation; anti-allergic; anti-asthmatic; asthma; anaphylaxis; dermatitis.
                                                                                                                                                                                                                   The present invention describes immunoglobulin R (IgB)-CH3 domain antigenic peptides (I). (I) have anti-allergic, anti-anaphylactic and anti-asthmatic properties. (I) induces polyclonal antibodies specific at target effector site on the epsilon-heavy chain of IgB, and so
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Canis sp.
Synthetic
 Sequence
                                                                                                                                                                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                                                                                                       New antigenic peptide from the CH3 for immunization against allergy.
                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-160578/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9967293-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Optimised IgB-CH3 domain antigen peptide for dog
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                               (UNBI-) UNITED BIOMEDICAL INC
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                                                                                                                                                                                                                                                                                            Page 99; 155pp; English.
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100.0%;
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Mismatches 0;
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RESULT 7
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Matches
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                                                                                                                                                                                                                                                                  AAW24098-106 are peptide fragments containing at least 5 continuous acids of the partial canine immunoglobulin E (IgB) protein shown in AAW24097. The peptides are used for the preparation of anti-canine I antibody. The anti-canine IgE antibody can be used for the diagnosis canine allergies
                                                                                                                                                                                                                                                                                                                                                  Canine immunoglobulin B peptide fragment and related DNA - useful for preparation of anti-canine immunoglobulin B antibody.
                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canine immunoglobin B peptide
                           Canine; allergy; antibody 15A.2;
epitope; prophylaxis; treatment;
                                                                                                 AAY50896;
                                                                                                                     AAY50896 standard; peptide;
                                                                                                                                                                                                                                                                                                                               Claim 2; Page 9; 12pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW24103 standard;
                                                          Antibody 15A.2 green monkey IgE binding epitope 1.
                                                                              24-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                 22-DEC-1995;
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                                                                                                                                                                     KDIVRSIAKAPGKRAP
                                                                                                                                                                                                          58.1%; S ilarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
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                              IgB; B cell; mast
mimotope.
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3. 6.1e-05;
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                                        cell; anti-allergy;
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Synthetic

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RESULT 8
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Matches 12
    Claim
                                                       New isolated polypeptides and encoding polymucleotides, useful for preventing and/or treating allergic diseases and allergy symptoms
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                                                  mammals.
                                                                                                                                         WPI; 2004-534123/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel binding protein which specifically binds to native canine free or B-cell bound IgE, and which doesn't bind to IgE when the IgE is bound to mast cells. The peptide products of the invention have anti-allergic activity: The antibodies bind to defined epitopes on free or B-cell bound IgE molecules which have an important role in allergic reaction. The specific binding proteins are used to produce a pharmaceutical composition, preferably with a diluent, which can be used for prophylaxis or treatment of canine allergy. AAY50876-y50900 represent peptide mimotopes used in the method of the invention
                                                                                                                                                                                                                                                                           20-DEC-2002; 2002US-00325375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        allergy; allergic symptom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide capable of preventing IgE binding
                                                                                                                                                                                                                                                                                                                      19-DEC-2003; 2003WO-US040711.
                                                                                                                                                                                                                                                                                                                                                                         15-JUL-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                    WO2004058799-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ76005 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Binding proteins used for treatment or prophylaxis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-040833/04.
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30-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lawton R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1999;
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                                                                                                                                                                                  Lawton R;
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50pp; English
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Query Match
Best Local Similarity
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                   This invention describes a novel binding protein which specifically binds to native canine free or B-cell bound IgE, and which doesn't bind to IgE when the IgE is bound to mast cells. The pertide products of the invention have anti-allergic activity. The antibodies bind to defined epitopes on free or B-cell bound IgE molecules which have an important role in allergic reaction. The specific binding proteins are used to produce a pharmaceutical composition, preferably with a diluent, which can be used for prophylaxis or treatment of canine allergy. ANY50876-Y50900 represent peptide mimotopes used in the method of the invention
Sequence
                                                                                                                                                           Disclosure; Fig 7; 30pp; English.
                                                                                                                                                                                   Binding proteins used for treatment or prophylaxis of
                                                                                                                                                                                                                 WPI; 2000-040833/04.
                                                                                                                                                                                                                                          Lawton R,
                                                                                                                                                                                                                                                                                               09-APR-1998;
30-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Canine; allergy; antibody 15A.2;
epitope; prophylaxis; treatment;
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                                                                                                                                                                                                                                                                                                                                                                17-NOV-1999.
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99US-00281760.
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Pred. No. 0.00062;
2; Mismatches 1;
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mimotope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to isolated polypeptides capable of inhibiting the binding of Igs to a high affinity receptor. These are useful for the prevention and/or treatment of allergic diseases and allergy symptoms in mammals. The present sequence is a polypeptide of the presen
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                                                                                                                                                                                                                                                                                                                 Immunoglobulin B; IgB; epsilon heavy chain; antigenic; antigen; immunogenic; immunosetimulatory; carrier protein; helper T cell epitope; antibody; allergy; allergic disease; immunisation; anti-allergic; anti-anaphylactic; anti-asthmatic; asthma; anaphylaxis; dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Optimised IgE-CH3 domain antigen peptide for rat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide capable of preventing IgE binding
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                                          29-DEC-1999
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73.3%;
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Pred. No. 0.0018;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local (
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                                                                                                                                                                                                                               Canine; allergy; antibody 15A.2; epitope; prophylaxis; treatment;
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                               Lawton R,
                                                                                    09-APR-1998;
30-MAR-1999;
                                                                                                                            09-APR-1999;
                                                                                                                                                                               BP957111-A2
                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                          Antibody 15A.2 swine IgB binding epitope 1.
                                                                                                                                                                                                                                                                                                                                 AAY50898;
                                                                                                                                                                                                                                                                                                                                                           AAY50898 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 99; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-160578/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                               8
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                               Mermer B,
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                                                                                                                                                                                                                                                                                                       (first entry)
                                                         LAB INC.
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99US-00281760
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WPI; 2000-040833/04.

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Best Local (
                                                                                    Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel binding protein which specifically binds to native canine free or B-cell bound IgE, and which doesn't bind to IgE when the IgE is bound to mast cells. The peptide products of the invention have anti-allergic activity. The antibodies bind to defined epitopes on free or B-cell bound IgE molecules which have an important role in allergic reaction. The specific binding proteins are used to produce a pharmaceutical composition, preferably with a diluent, which can be used for prophylaxis or treatment of canine allergy. AAY50876-
                                                                                                                         The present invention relates to isolated polypeptides capable of inhibiting the binding of IgE to a high affinity receptor. These ar useful for the prevention and/or treatment of allergic diseases and allergy symptoms in mammals. The present sequence is a polypeptide of the present sequence is a polypeptide of the present sequence.
                                                                                                                                                                                                      Claim 5; SEQ ID
                                                                                                                                                                                                                                           New isolated polypeptides and encoding polynucleotides, useful for preventing and/or treating allergic diseases and allergy symptoms
                                                                                                                                                                                                                                                                                            WPI; 2004-534123/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004058799-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                allergy; allergic symptom; IgE binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-OCT-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADQ76007;
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                                                                                                                                                                                                                                                                                                                                                       (IDEX-) IDEXX LAB INC
                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-2002; 2002US-00325375
                                                                                                                                                                                                                                                                                                                                                                                                              19-DEC-2003; 2003WO-US040711.
                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide capable of preventing IgE binding to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 AA;
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                          12;
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CRVTHPHLPKDIVRSI 16
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75.0%;
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75.0%;
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                          Score 65; DB Pred. No. 0.00
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0.0075;
                                      DB 8;
0.0075;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          high affinity receptor
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AAY91212;

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AAY68602
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                                                                                                                                                               Query Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                The specification describes peptide immunogens comprising a synthetic helper T cell (Th) epitope and a target antigen, luteinising hormone-releasing hormone (LHRH). The peptide immunogens cause induction of a specific immune response to LHRH which is involved in regulation of spermatogenesis, ovulation, oestrus, sexual development and secretion of sex hormones. Provision of a promiscuous T helper epitope (which is functional in genetically diverse subjects) provides optimum immunogenicity to the B cell epitopes of the target antigen and thus high antibody titres against the target antigen. The peptide immunogens of the invention are used to vaccinate against mammalian LHRH, for use as (reversible) contraceptive, control of hormone-dependent tumours (cancer of prostate or breast, also endometriosis); to prevent boar taint (and improve meat quality) and for immunocastration. The present sequence
                               AAY91212
                                                                                                                                                                                                                           Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide immunogen containing luteinizing hormone-releasing antigen site and helper T cell epitope, for e.g. contraception treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Helper T cell epitope; peptide immunogen; LHRH;
luteinising hormone-releasing hormone; spermatogenesis; ovulation;
oestrus; sexual development; sex hormone; promiscuous T helper epitope;
vaccine; contraceptive; hormone-dependent tumour; prostate cancer;
breast cancer; endometriosis; boar taint; meat quality; immunocastration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 92; 102pp; English
                                                                                                                                                                                                                                                       appears in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-160562/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUN-1998;
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                            standard; peptide; 25
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The invention relates to novel promiscuous T helper cell epitopes (Th), calling immunogenic peptides comprising the Th epitopes of the invention call and immunogens generally against plasmatch to induce a T halper cell response, containing them, are used to induce a T halper cell response, containing them, are used to induce a T halper cell response, peptide immunogens containing them, are used to induce a T halper cell response, containing them, are used to induce a T halper cell response, petitically against any pathogen, containing them or through antique. The Th epitopes and peptide immunogens may be used for prevention and/or treatment of peptide immunogens may be used for prevention and/or treatment of peptide immunogens may be used for prevention and/or treatment of peptide immunogens may be used for prevention of lutainising hormone contains the promotent of antibition of the action of lutainising hormone contains the promotent of antibition of the action of lutainising hormone contains the promotent of the period into an immunogen improves capacity to induce a strong T helper cell-mediated immune response, resulting in production of antibodies against a target antigen. The can replace carrier contains and pathogen-derived T helper epitopes. Sequence ANY31121 periods and ANY31221 periods contains and pathogen-derived T helper epitopes from the measles virus F (MVF) corrects antigent peptides comprising an tirk sequence sequence ANY31121. ANY31226 and ANY31245-Y91246 periods antigent peptides antigent peptide corrects antigent peptide corrects antigent peptide comprising somatostatin, and CC ANY3120-Y9101 are antigent peptides. ANY311370 is somatostatin, and CC ANY3120-Y9101 are matigent peptides. ANY31270 is somatostatin, and antigent of alman 1gg (immunogens may be used to promote growth in custok. ANY3121 are MVF in epitope CDA CDR2 antigent peptides which may be used in the custok. ANY3121 are mVF in epitope and period derived from foot and mouth custome of a luman 1gg (immunogens may be used in the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigenic site, for immunization or human immune deficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New artificial T helper cell epitope and derived immunogens with target antigenic site, for immunization against e.g. malaria, arteriosclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Promiscuous T-cell epitope; measles virus P protein; MVP; hepatitis B virus surface antigen; HBV; humungenic; B-cell epitope; luteinising hormone releasing hormone; LHRH; contraceptive; anticancer; somatostatin; growth promotion; CD4 receptor; HIV-1; antiviral; PMDV; foot and mouth disease virus; immunoglobulin B; IGB; anti-allergic; Plasmodium falciparum; circumsporzoite; antimalarial; CETP; cholesteryl ester transport protein; anti-arteriosclerotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 40; 129pp; English.
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Matches
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                                                                                                                                                                                                                          Sequence 25 AA,
                                                                                                                                        Local Similarity
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B
                                                    2 RYTHPHLPKDIVRSIAK 18
RVTHPHLPRALMRSTTK 24
                                                                                                           Conservative
                                                                                                                                  47.18;
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Search completed: February 22, 2005, 07:16:58 Job time : 163 secs

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Post-processing: Minimum Match 0%
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Perfect score:
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1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

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10: /cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

13: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

14: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

15: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

16: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

16: /cgn2_6/ptodata/2/pubpaa/US10D_PUBCOMB.pep:*

17: /cgn2_6/ptodata/2/pubpaa/US10NEW_PUB.pep:*

18: /cgn2_6/ptodata/2/pubpaa/US10NEW_PUB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US10B_PUB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US10B_PUB.pep:*

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63.914 Million cell updates/sec
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US-10-325-375A-3
US-10-325-375A-6
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US-10-630-074-96
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US-10-325-375A-5
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US-10-325-375A-5
US-10-344-43-107
US-10-344-43-107
US-10-344-43-197
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Sequence 50, Appli
Sequence 107, Appli
Sequence 20, Appli
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Sequence 27, Appli
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Patent No. US20020064525A1

GENERAL INFORMATION:
APPLICANT: MOTSEY, et al.
TITLE OF INVENTION: Anti-IgE Vaccines
FILE REFERENCE: PC10761A
CURRENT APPLICATION NUMBER: US/09/938,700
CURRENT ETLING DATE: 2001-08-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 25
TYPE: PRI
TYPE: PRI
TYPE: PRI
TYPE: PRI
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US-10-325-375A-2
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US-10-362-527-62	-10-304-443-	-10-322-210	-10-415-389-	-10-415-389-	-527-	-10-362-527-	-10-304-443-	-10-304-443-	-10-415-389-	-10-415-389-	US-10-304-443-119	US-10-304-443-96	9-5	0-415	10-415-389-	-10-362-527-	-10-362-527-	-10-304-443-	-10-304-	-10-415-389-	-10-415-389-	-10-415-389-	-10-415-389-	-10-362-527-3	62-527-	-10-304-443-1	-10-304-443-	-10-304-443-	US-10-304-443-98	US-10-415-389-33	US-10-415-389-10
Sequence 62, Appl	equence 3,		e 27	e 21	e 32	e 31	e 11	e 10	32	е 9,	115	O	33,	29,	e 23,	ø	316,	116	æ	e 34,	e 28,	e 22,	e 11,	e 321	e 315,	æ	e 115	e 109	e 98,	e 33,	Ф

ALIGNMENTS

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Sequence 2, Application US/10325375A
Publication No. US20030229021A1
GENERAL INFORMATION:
APPLICANT: IDEXX Laboratories, Inc.
APPLICANT: Krah, Eugene R.
APPLICANT: Krah, Eugene R.
APPLICANT: Lawron, Robert
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of IgE to a High
                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 136; DB 9; Best Local Similarity 100.0%; Pred. No. 4.6e-12; Matches 25; Conservative 0; Mismatches 0;
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US-10-325-375A-3
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US-10-325-375A-4
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APPLICANT: Krah, Eugene R.
APPLICANT: Lawton, Robert
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of IgE to a High
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of IgE to a High
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of IgE to a High
TITLE REFERENCE: MHBH-01-672-8
CURRENT APPLICATION INVESER: US/10/325,375A
CURRENT FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn version 3.2
SEQ ID NO 4
LENGTH: 17
                                                                                                                                                                                                                            Sequence 3, Application US/10325375A Publication No. US20030229021A1 GENERAL INFORMATION:
SOFTWARE: PatentIn
SEQ ID NO 3
LENGTH: 17
                                                APPLICANT: IDEXX Laboratories, Inc.
APPLICANT: Lawton, Eugene R.
APPLICANT: Lawton, Robert
TITLE OF INVENTION: Methods and Compositions for
FILE REFERENCE: MBHB-01-672-E
CURRENT APPLICATION NUMBER: US/10/325,375A
CURRENT FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 14
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Best Local (
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SEQ ID NO 2
LENGTH: 17
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Affinity Receptor FILE REFERENCE: MBHB-01-672-E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Isolated polypeptide that binds to IgE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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100.0%; Pred. No. 1.7
tive 0; Mismatches
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Pred. No. 0.00
2; Mismatches
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hes 0;
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                                                                                                                                         Inhibiting Binding of IgE to a High
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Sequence 96, Application US/10630070
Publication No. US20050025781A1
GENERAL INFORMATION:
APPLICANT: Milch, David R.
APPLICANT: Milch, David R.
TITLE OF INVENTION: Rodent Hepatitis B Virus Co:
TITLE OF INVENTION: Methods of Use Thereof
FILE REFERENCE: VACCINE-07083
CURRENT APPLICATION NUMBER: US/10/630,070
CURRENT FILING DATE: 2003-07-30
                                                                      ; OTHER INFORMATION: Synthetic US-10-630-070-96
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US-10-325-375A-6
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                                                                                                                                 NUMBER OF SEQ ID NOS: 101
SOFTWARE: PatentIn version 3.2
SEQ ID NO 96
LENGTH: 23
TYPE: PRT
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Best Local S
Matches 12
        Matches
                    Query Match
Best Local Similarity
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SEQ ID NO 6
LENCTH: 17
TYPE: PRT
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: IDEXX Laboratories, Inc.
APPLICANT: Krah, Eugene R.
APPLICANT: Lawton, Robert
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of
TITLE OF INVENTION: Affinity Receptor
FILE REFERENCE: MBHB-01-672-B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/325,375A
CURRENT FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 14
                                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Isolated polypeptide that binds to IgE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial FEATURE: OTHER INFORMATION: Isolated polypeptide that binds to IgE.
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      Conservative
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llarity 75.0%; Pred. No. 0.029;
Conservative 1; Mismatches
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                  47.18;
64.78;
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 Score 64; DB
Pred. No. 0.05
3; Mismatches
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Pred. No. 0.008;
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                  0.056;
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                                Length 23;
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RVTHPHLPKDIVRSIAK 18

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US-10-304-443-107
US-10-304-443-107

### Sequence 107, Application US/10304443

### Publication No. US20030170229A1

#### GENERAL INFORMATION:
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Best Local
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LENGTH: 17
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Best Local
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CURRENT FILING DATE: 2003-07-30
NUMBER OF SEQ ID NOS: 101
SOFTWARE: PatentIn version 3.2
SEQ ID NO 96
LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 96, Application US/10630074
Publication No. US20050025782A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: IDEXX Laboratories, Inc.
APPLICANT: Krah, Eugene R.
APPLICANT: Lawton, Robert
TITLE OF INVENTION: Methods and Compositions for Inhibiting Binding of IgE to a High
TITLE OF INVENTION: Affinity Receptor
FILE REFERENCE: MBHB-01-672-B
CURRENT FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Milich, David R.
APPLICANT: Millaud, Jean-Noel
APPLICANT: Billaud, Jean-Noel
TITLE OP INVENTION: Human Hepatitis B Virus Core Proteins as
TITLE OP INVENTION: Methods of Use Thereof
FILE REFERENCE: VACCINE-07971
APPLICANT: Smithkline Beecham Biologicals s.a.
                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Isolated polypeptide that binds to IgE.
                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial
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75.0%; Pred. No. 0.
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RESULT 11 US-10-415-389-20

Sequence 20, Application US/10415389
Publication No. US20040115220A1
GENERAL INFORMATION:
APPLICANT: De Basols, Carlota Vinals Y
TITLE OF INVENTION: Vaccine

PILE REFERENCE: B45251
CURRENT APPLICATION NUMBER: US/10/415,389
CURRENT FILING DATE: 2003-04-25
PRIOR APPLICATION NUMBER: PCT/EP01/12932
PRIOR FILING DATE: 2001-10-24

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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 107
LENGTH: 14
                                                                           Query Match
Best Local Similarity
Matches 9; Conservat
                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 328
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 313
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/362,527
CURRENT FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: PCT/EP01/09576
PRIOR FILING DATE: 2001-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Friede, Martin
APPLICANT: Mason, Sean
APPLICANT: Turnell, William Gordon
APPLICANT: Vinals Y De Bassols, Carlota
APPLICANT: Vinals Y De Carlota
APPLICANT: NURSITION: Vaccine Immunogens Compx
TITLE OF INVENTION: and Use Thereof in the
FILE REFERENCE: B45236
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TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45173CIP
CURRENT APPLICATION NUMBER: US/10/304,443
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US/09/698,906A
PRIOR APPLICATION NUMBER: US/09/698,906A
PRIOR FILING DATE: 2001-02-20
                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: GB 0020717.5
PRIOR FILING DATE: 2000-08-22
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                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                               OTHER INFORMATION: Artificial variant of Homo sapiens IgB peptide
                                                                                                                                                                                                                                                         ENGTH: 14
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                                   СКУТНРИГЬКОІ 13
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CRVTHPHLPRALM 13
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                                                                           Conservative
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                                                                                              44.1%;
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                                                                           3; Mismatches
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                                                                                                Score 60; DB 1
Pred. No. 0.12;
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Pred. No. 0.12;
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US-10-304-443-120
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; LOCATION: (2)
; OTHER INFORMATION: Where Xaa represents any 1 of 20 naturally occurring amino acids
US-10-304-443-97
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                                                                                                                                                               Sequence 120, Application US/10304443
Publication No. US20030170229A1
GENERAL INFORMATION:
APPLICANT: Smithkline Beecham Biologicals
APPLICANT: Peptide Therapeutics Ltd.
TITLE OF INVENTION: Vaccine
SEQ ID NO 120
LENGTH: 16
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Best Local (
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CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US/09/698,906A
PRIOR FILING DATE: 2001-02-20
NUMBER OF SEQ ID NOS: 121
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Smithkline Beecham Biologicals s.a.
APPLICANT: Peptide Therapeutics Ltd.
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45173CIP
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PRIOR FILING DATE: 2000-10-27
NUMBER OF SEQ ID NOS: 34
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 20
LENGTH: 14
TYPE: PRT
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                                                                    CURRENT APPLICATION NUMBER: US/10/304,443
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US/09/698,906A
PRIOR FILING DATE: 2001-02-20
                                 NUMBER OF SEQ ID NOS: 121
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                     FILE REFERENCE: B45173CIP
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ORGANISM: Human peptide sequence
FEATURE:
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Pred. No. 0.14
3; Mismatches
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RESULT 15
US-10-415-389-33
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                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 33
                                                                                                                                                                                                                                                         Sequence 33, Application US/10415389
Publication No. US20040115220A1
GENERAL INFORMATION:
APPLICANT: De Basols, Carlota Vinals Y
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10
LENGTH: 16
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Matches
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Best Local Similarity
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                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/415,389
CURRENT FILING DATE: 2003-04-25
PRIOR APPLICATION NUMBER: PCT/EP01/12932
PRIOR FILING DATE: 2001-10-24
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                                                                                                                                 NUMBER OF SEQ ID NOS:
                                                                                                                                                PRIOR APPLICATION NUMBER: GB 0026334.3 PRIOR FILING DATE: 2000-10-27
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CURRENT FILING DATE: 2003-04-25
PRIOR APPLICATION NUMBER: PCT/EP01/12932
PRIOR FILING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: GB 0026334.3
PRIOR APPLICATION NUMBER: GB 0026334.3
PRIOR FILING DATE: 2000-10-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: De Basols, Carlota Vinals Y TITLE OF INVENTION: Vaccine FILE REFERENCE: B45251
OTHER INFORMATION: Human IgE peptide mimotope
                      ORGANISM: Artificial Sequence FEATURE:
                                                           TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 34
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NAME/KEY: VARIANT
LOCATION: (1)...(16)
OTHER INFORMATION: Xaa = Any Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Human IGE peptide mimotope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Human peptide sequence
                                                                            ENGTH:
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Similarity 71.4%;
10; Conservative
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10. US20040115220A1
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Pred. No. 0.14
3; Mismatches
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Pred. No. 0.14;
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Query Match
Query Match
Best Local Similarity 71.4%; Pred. No. 0.14;
Matches 10; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 2 RVTHPHLPKDIVRS 15
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Db 3 RVTHPHLPRALMRS 16

Search completed: February 22, 2005, 07:32:24

Job time: 128 secs
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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    35 55
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seq length: 25
     Query
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1: /cgn2_6/ptcdata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptcdata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptcdata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptcdata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptcdata/1/iaa/FCTUS_COMB.pep:*
6: /cgn2_6/ptcdata/1/iaa/backfiles1.pep:*
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42.414 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd
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US-09-281-7608-27
US-09-281-7608-30
US-09-201-623C-3
US-09-701-623C-5
US-09-701-623C-5
US-09-701-623C-5
US-09-701-623C-5
US-09-701-623C-5
US-09-701-623C-5
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US-09-701-623C-5
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US-09-281-7608-29
US-09-281-7608-31
US-09-281-7608-18
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US-09-281-7608-18
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US-09-281-7608-18
US-09-281-7608-19
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US-09-701-623C-6
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Sequence 26, Appl
Sequence 6, Appli
Sequence 27, Appl
Sequence 97, Appl
Sequence 95, Appl
Sequence 95, Appl
Sequence 95, Appl
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Sequence 97, Appl
Sequence 98, Appl
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Sequence 97, Appl
Sequence 97, Appl
Sequence 98, Appl
Sequence 19, Appl
Sequence 29, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 12, Appl
Sequence 13, Appl
Sequence 14, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 12, Appl
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Sequence 6, Application US/09701623C

Patent No. 6811782

GENERAL INFORMATION:
APPLICANT: Wang Ph.D., Chang Yi
ITILE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF INTERGY
PILE REFERENCE: 11514153US1

CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: PCT/US99/13959

PRIOR RILING DATE: 1999-06-21
PRIOR FILING DATE: 1999-06-21
PRIOR FILING DATE: 1998-06-20

NUMBER OF SEQ ID NOS: 91
SOPTWARE: PATENTIN NOW: 2.1
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US-09-043-222C-105 US-08-478-684G-12 US-09-046-985-4 US-09-474-743-4 US-08-851-843A-208 US-08-854-050-208 US-08-974-549A-327 US-08-973-323-208 US-09-402-181B-327 US-09-4731-456-327 US-08-480-190-80 US-08-480-190-80 US-08-480-190-80 US-08-480-379-80 US-08-480-379-80 US-08-471-255A-80 US-08-851-153A-5 US-08-861-153A-5 US-08-861-153A-5	4	4	ហ	4	4	N	N	N	4	4	w	w	w	w	w	w	۵	4
	US-09-493-795B-79	US-08-861-153A-5	PCT-US93-07545-80	US-08-077-255A-80	US-08-475-399A-80	US-08-488-379-80	US-08-480-190-80	US-08-323-686-15	US-09-721-456-327	US-09-402-181B-327	US-09-430-323-208	US-08-854-050-208	US-08-974-549A-327	US-08-851-843A-208	-09	-09	US-08-478-684G-12	
	79, Appl	5, Appli	80, Appl	•	80, Appl	•	•	•	327, App	327, App	•	•	327, App	•	4, Appli	4, Appli	12, App1	•

ALIGNMENTS

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APPLICANT: Lawton, Robert
APPLICANT: Mermer, Brion
APPLICANT: Mermer, Brion
APPLICANT: Mermer, Brion
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
CURRENT APPLICATION NUMBER: US/09/281,760B
CURRENT PILING DATE: 199-03-30
FRIOR APPLICATION NUMBER: 09/058,331
FRIOR APPLICATION NUMBER: 09/058,331
FRIOR FILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
SOPTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 26
SEQ ID NO 26
SEQ TO 26
ORGANISM: Canis familiaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 26, Application US/09281760E Patent No. 6734287
GENERAL INFORMATION:
                                                                                                   Query Match
Best Local Similarity
                     1 CRVTHPHLPKDIVRSI 16
CRVTHPHLPKDIVRSI
                                                                                 Conservative
                                                                         65.4%; or
100.0%; Pr
  17
                                                                                                     Score 89; DB 4; I; Pred. No. 2.1e-07;
                                                                                 Mismatches
                                                                                                                          Length 17;
                                                                                   0; Indels
                                                                                   0,
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Gaps

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APPLICANT: Francoeur, Greg
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
CURRENT APPLICATION NUMBER: US/09/281,760E
CURRENT FILING DATE: 1999-03-30
FRIOR APPLICATION NUMBER: 09/058,331
FRIOR APPLICATION NUMBER: 09/058,331
PRIOR FILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 28
LENCTH: 17
                                                       ; ORGANISM: Homo sapiens US-09-281-760E-27
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; ORGANISM: Cercopithecus aethiops
US-09-281-760E-28
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                                                                                                                               NUMBER OF SEQ ID NOS: 39
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 27
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 27, Application US/09281760E Patent No. 6734287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 52.9%;
Best Local Similarity 80.0%;
Matches 12; Conservative
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GENERAL INFORMATION:
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Best Local :
                                                                                                                                                                                                                                                                  APPLICANT: Lawton, Robert
APPLICANT: Mermer, Brion
APPLICANT: Francoeur, Greg
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
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CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 09/058,331
PRIOR FILING DATE: 1998-04-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lawton, Robert APPLICANT: Mermer, Brion APPLICANT: Francoeur, Gr
                                                                                                                 LENGTH: 17
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ORGANISM: Artificial Sequence
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 50.7%;
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 Score
Pred.
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No.
DB 4;
0.00029;
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               Length 17;
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CURRENT APPLICATION NUMBER: US/09/281,760B
CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 09/058,331
PRIOR FILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
SOFTMARE: PASTSEQ for Windows Version 3.0
SEQ ID NO 30
LENGTH: 17
TYPE: PRT
ORGANISM: Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Wang Ph.D., Chang Yi
APPLICANT: Wang Ph.D., Chang Yi
ITITLE OF INVENTION: ALLERGY
FILE REFERENCE: 11514153US1
CURRENT APPLICATION NUMBER: US/09/701,623C
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: PCT/US99/13959
PRIOR FILING DATE: 1999-06-21
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR APPLICATION NUMBER: 09/100,287
PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
SOPTWARE: Patentin Ver. 2.1
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US-09-281-760E-30
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US-09-701-623C-7
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                                                                  Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Lawton, Robert
APPLICANT: Mermer, Brinn
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30, Application US/09281760E Patent No. 6734287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/09701623C
Patent No. 6811782
                                                                                                                                                                                                                                                                                                                    APPLICANT: Mermer, Brion
APPLICANT: Francoeur, Greg
TITLE OF INVENTION: Specific Binding Protein for Treating
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-125A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 25
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                  Local Similarity hes 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CRVTHPHLPKDIVRS
                              CRVTHPHLPKDIVRSI 16
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CNVTHPDLPKPILRSI 17
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                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                 47.8%; Score 65; DB 4; Length 17; 75.0%; Pred. No. 0.0013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50.0%; Score 68; DB 4; 76.5%; Pred. No. 0.00066;
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                                                                Mismatches
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                                                                Gaps
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; Sequence 95, Application US/09303323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
COUNTRY: USA
ZIP: 10154-0054
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC Windows
SOFTWARE: Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/303,323
FILING DATE: 30-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Wang, (
                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Wang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AGENT INFORMATION:
NAME: MARIA H. Lin
REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 1151
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212,758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 212-751-6849
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wang, Chang Yi
TITLE OF INVENTION: NOVEL LHRH
TITLE OF INVENTION: IMMUNOGENS
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC Windows
SOFTWARE: Word 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/100,414B
FILING DATE: 20-JUNB-1998
CLASSIFICATION: 424
                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Morgan &
                                                                                                                                                                                                                                                                                                          APPLICANT: Wang, Chang Yi
TITLE OF INVENTION: NOVEL LHRH PEPTIDE
TITLE OF INVENTION: IMMUNOGENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 10154-0054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Morgan & Finnegan, L.L.P. STREET: 345 Park Avenue
                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 25 amino acids
                                                                                                                                                                                            TATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RVTHPHLPRALMRSTTK 24
                                                                                                                                                                                                                                 B: Morgan & Finnegan, L.L.P. 345 Park Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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Pred. No. 0.0028;
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US-09-770-014-95
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                                                US-09-770-014-95
                                                                               TELEPHONE: 212-758-4800
TELEPAX: 212-751-6849
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Best Local Similarity 64.7%;
Matches 11; Conservative
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 212-751-6849
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/100,414
FILING DATE: 20-JUNE-1998
ATTORNEY/AGENT INFORMATION:
NAME: Maria H. Lin
REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 1151-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOV.
TITLE OF INVENTION: IMM
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                        SUPTWARE: WORD 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/770,014
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
LENGTH: 25 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION
TELEPHONE: 212-758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 20-JUNE-199
ATTORNEY/AGENT INFORMATION:
NAME: Maria H. Lin
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                             TOPOLOGY: 11
                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 212-758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: Z12-751-6849
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 29,323
                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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STREET: 345 Park Avenue
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                                                                                                                                                                                                                                                                                                                                                                                          IBM PC compatible SYSTEM: PC Windows
                                                           peptide
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 47.1%;
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   Score 64; DB 4;
Pred. No. 0.0028;
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RESULT 11

US-09-701-623C-5

Sequence 5, Application US/09701623C

Patent No. 6811782

GENERAL INFORMATION:

APPLICANT: Wang Ph.D., Chang Yi

TITLE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR TITLE REFERENCE: 11514153US1

CURRENT APPLICATION NUMBER: US/09/701,623C

CURRENT FILING DATE: 2000-12-01

PRIOR APPLICATION NUMBER: PCT/US99/13959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
US-09-701-588C-92
US-09-701-588C-92
; Sequence 92, Application US/09701588C
; Patent No. 6713301
; GENERAL INFORMATION:
; APPLICANT: UNITED BIOMEDICAL INC., ET AL.
; APPLICANT: UNITED BIOMEDICAL T HELPER CELL
. TITLE OF INVENTION: ARTIFICIAL T HELPER CELL
. EPITOPES AS IMMUNE STIMULATORS FOR SYNTHETIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 1151
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-758-4800
TELEPAX: 212-751-6849
INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC WINDOWS
SOFTWARE: WOX 97
CURRENT APPLICATION NUMBER: US/09/701,588C
APPLICATION NUMBER: US/09/701,588C
FILING DATE: 29-No. 6713301-2002
CIASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/100,414
FILING DATE: 20-UNE-1998
ATTORNEY/AGENT INFORMATION:
NAME: MATIE H. Lin
                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
nes 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Morgan & Finnegan, L.L.P.
STREET: 345 Park Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 92:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10154-0054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                           47.1%; Score 64; DB 4; Length 25; ilarity 64.7%; Pred. No. 0.0028; Conservative 3; Mismatches 3; Indels
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APPLICANT: Wang Ph.D., Chang Yi

APPLICANT: Wang Ph.D., Chang Yi

ITILE OF INVENTION: PEPTIDE COMPOSITION AS IMMUNOGEN FOR THE TREATMENT OF

TITLE OF INVENTION: ALLERGY

FILE REFERENCE: 11514153US1

CURRENT APPLICATION NUMBER: US/09/701,623C

CURRENT FILING DATE: 2000-12-01

PRIOR APPLICATION NUMBER: PCT/US99/13959

PRIOR APPLICATION NUMBER: 09/100,287

PRIOR APPLICATION NUMBER: 09/100,287

PRIOR APPLICATION NUMBER: 1998-06-21

SOFTWARE: PATENT 1998-06-20

NUMBER OF SEQ ID NOS: 91

SOFTWARE: PATENTIN Ver. 2.1
                       GENERAL INFORMATION:
APPLICANT: Lawton, Robert
APPLICANT: Mermer, Brion
APPLICANT: Mermer, Brion
APPLICANT: Francoeur, Greg
TITLE OF INVENTION: Specific Binding Protein for
TITLE OF INVENTION: Canine Allergy
FILE REFERENCE: 01-1275A
CURRENT APPLICATION NUMBER: US/09/281,7608
CURRENT APPLICATION NUMBER: 09/08-331
PRIOR APPLICATION NUMBER: 09/08,331
PRIOR PILING DATE: 1998-04-09
NUMBER OF SEQ ID NOS: 39
NUMBER OF SEQ ID NOS: 39
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PRIOR APPLICATION NUMBER: 09/10
PRIOR FILING DATE: 1998-06-20
NUMBER OF SEQ ID NOS: 91
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: synthetic US-09-701-623C-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
US-09-701-623C-84
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                                                                                                                                                                                                                                                                                                                         US-09-281-760E-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-701-623C-84
                                                                                                                                                                                                                                                       Sequence 29, Application US/09281760E Patent No. 6734287
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 84,
Patent No. 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
          SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 25
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Pred. No.
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0.004;
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; SEQ ID NO 29
; LENGTH: 17
; TYPE: PRT
; ORGANISM: Felis catus
US-09-281-760E-29
                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Watches 9; Conserve
RESULT 15

US-08-232-539D-20

; Sequence 20, Application US/08232539D

; Patent No. 5965709
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MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Winpatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,539D
FILING DATE: 21-Apr-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/178583
FILING DATE: 07-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/744768
FILING DATE: 07-JAN-1994
PRIOR APPLICATION NUMBER: 09/1744768
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: SYODOGA, CTRAIG G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: 39,044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19, Application US/08232539D Patent No. 5965709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 44.9%;
Best Local Similarity 75.0%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
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APPLICANT: Jardieu, Paula M.
TITLE OF INVENTION: 19E Antagonists
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1 DNA Way
CITY: South San Francisco
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                                                                                                                                                                                            1 CRVTHPHLPKDIV 13
|||||||:::
10 CRVTHPHLPRALM 22
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                                                                                                                                                                                                                                                                                                                                                          44.1%;
69.2%;
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Pred. No. 0.01;
3; Mismatches
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Search completed: February 22, 2005, 07:29:40 Job time : 45 Becs

Gaps

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COMPUTER READABLE FORM,

MEDIUM TYPE: J.5 inch, 1.44 Mb floppy diek

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/232,539D

FILING DATE: 21-Apr-1994

CLASSIFICATION NUMBER: 08/178583

FILING DATE: 07-JAN-1994

PRIOR APPLICATION NUMBER: 07/744768

FILING DATE: 14-AUG-1991

APPLICATION NUMBER: 07/744768

FILING DATE: 14-AUG-1991

APPLICATION NUMBER: 07/744768

FILING DATE: 14-AUG-1991

APPLICATION NUMBER: 99/14768

FILING DATE: 14-AUG-1994

APPLICATION NUMBER: 99/14768

APPLICATI
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Matches
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APPLICANT: Presta, Leonard G.
APPLICANT: Jardieu, Paula M.
TITLE OF INVENTION: 1GB Antagonists
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1 STREET: 1 CITY: South San Fra
                                                                                                                                                       y Match 44.1%;
Local Similarity 69.2%;
les 9; Conservative
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12 CRVTHPHLPRALM 24
                                                                               1 СКУТНРНГЬКОІ 13
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                                                                                                                                                               Score 60; DB 2;
Pred. No. 0.011;
3; Mismatches
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